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54 Plant cells resistant to glutamine synthetase inhibitors, made by genetic engineering.

57 The invention relates to a DNA fragment containing a determined gene, the expression of which inhibits the antibiotic and herbicidal effects of Bialaphos and related products.

It also relates to recombinant vectors, containing such DNA fragment, which enable this protective gene to be introduced and expressed into cells and plant cells.

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GENETICALLY ENGINEERED PLANT CELLS AND PLANTS EXHIBITING  
RESISTANCE TO GLUTAMINE SYNTHETASE INHIBITORS, DNA FRAG-  
MENTS AND RECOMBINANTS FOR USE IN THE PRODUCTION OF SAID  
5 CELLS AND PLANTS.

The invention relates to a process for protecting  
plant cells and plants against the action of glutamine  
synthetase inhibitors.

It also relates to applications of such process,  
10 particularly to the development of herbicide resistance  
into determined plants.

It relates further to non-biologically transformed  
plant cells and plants displaying resistance to glutamine  
synthetase inhibitors as well as to suitable DNA fragments  
15 and recombinants containing nucleotide sequences encoding  
resistance to glutamine synthetase inhibitors.

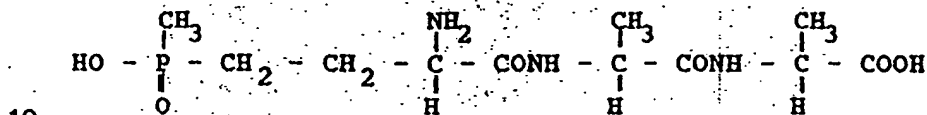
Glutamine synthetase (hereafter simply designated  
by GS) constitutes in most plants one of the essential  
enzymes for the development and life of plant cells. It is  
20 known that GS converts glutamate into glutamine. GS is  
involved in an efficient pathway (the only one known  
nowadays) in most plants for the detoxification of ammonia  
released by nitrate reduction, aminoacid degradation or  
photorespiration. Therefore potent inhibitors of GS are  
25 very toxic to plant cells. A particular class of  
herbicides has been developped, based on the toxic effect  
due to inhibit inhibition of GS in plants.

These herbicides comprise as active ingredient a  
GS inhibitor.

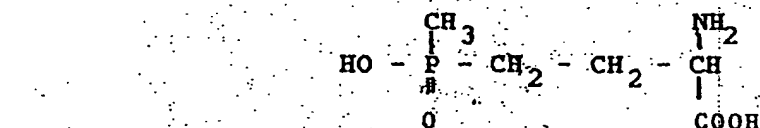
30 There are at least two possible ways which might  
lead to plants resistant to the inhibitors of the action  
of glutamine synthetase ; (1) by changing the target. It  
can be envisaged that mutations in the GS enzyme can lead  
to insensitivity towards the herbicide ; (2) by inactiva-  
35 tion of the herbicide. Breakdown or modification of the  
herbicide inside the plant could lead to resistance.

Bialaphos and phosphinothricin (hereafter simply designated by PPT) are two such inhibitors of the action of GS, (ref. 16, 17) and have been shown to possess excellent herbicidal properties (see more particularly ref. 2 as concerns Bialaphos).

Bialaphos has the following formula (I) :



PPT has the following formula (II) :



Thus the structural difference between PPT and Bialaphos resides in the absence of two alanine aminoacids in the case of PPT.

These two herbicides are non selective. They inhibit growth of all the different species of plants present on the soil, accordingly cause their total destruction.

Bialaphos was first disclosed as having antibiotic properties, which enabled it to be used as a pesticide or a fungicide. Bialaphos can be produced according to the process disclosed in united-states patent n° 3 832 394, assigned to MEIJI SEIKA KAISHA LTD., which patent is incorporated herein by reference. It comprises cultivating Streptomyces hygroscopicus, such as the strain available at the American Type Culture Collection, under the ATCC number 21,705, and recovering Bialaphos from its culture medium. However, other strains, such as Streptomyces viridochromogenes, also produce this compound (ref. 1).

Other tripeptide antibiotics which contain a PPT moiety are or might be discovered in nature as well, e.g. phosalacin (ref. 15).

PPT is also obtained by chemical synthesis and is commercially distributed by the industrial Company

HOESCHT.

5 A number of Streptomyces species have been disclosed which produce highly active antibiotics which are known to incapacitate procaryotic cell functions or enzymes. The Streptomyces species which produce these antibiotics would themselves be destroyed if they had not a self defence mechanism against these antibiotics. This  
10 self defence mechanism has been found in several instances to comprise an enzyme capable of inhibiting the antibiotic effect, thus of avoiding autotoxicity for the Streptomyces species concerned. This modification is generally reversed when the molecule is exported from the cell.

15 ~~The existence of a gene which encodes an enzyme able to modify the antibiotic so as to inhibit the antibiotic effect against the host has been demonstrated in several Streptomyces producing antibiotics, for example, in S. fradiae, S. azureus, S. vinaceus, S. erythreus, producing neomycin, thiostrepton, viomycin, and MLS (Macrolide Lincosamide Streptogramin) antibiotics respectively (ref. 4), (ref. 5), (ref. 6), (ref. 14 by CHATER et al., 1982 describes standard techniques which can be used for bringing these effects to light).~~

25 In accordance with the present invention, it has been found that Streptomyces hygroscopicus ATCC 21,705, also possesses a gene encoding an enzyme responsible of the inactivation of the antibiotic properties of Bialaphos. Experiments carried out by the applicants have  
30 lead to the isolation of such a gene and its use in a process for controlling the action of GS inhibitors, based on PPT or derived products.

35 An object of the invention is to provide a new process for controlling the action in plant cells and plants of GS inhibitors.

Another object of the invention is to provide DNA

fragments and DNA recombinants, particularly modified vectors containing said DNA fragments, which DNA fragments contain nucleotide sequences capable, when incorporated in plant cells and plants, to protect them against the action of GS inhibitors.

A further object of the invention is to provide non-biologically transformed plant cells and plants capable of neutralizing or inactivating GS inhibitors.

A further object of the invention is to provide a process for selectively protecting plant species against herbicides of a GS inhibitor type.

More specifically an object of the invention is to provide a DNA fragment transferable to plant cells- and to whole plants- capable of protecting them against the herbicidal effects of Bialaphos and of structurally analogous herbicides.

A further object of the invention is to provide plant cells resistant to the products of the class exemplified by Bialaphos, which products possess the PPT unit in their structure.

The process according to the invention for controlling the action in plant cells and plants of a GS inhibitor when contacted therewith, comprises providing said plants with a heterologous DNA fragment including a foreign nucleotide sequence, capable of being expressed in the form of a protein in said plant cells and plants, under condition such as to cause said heterologous DNA fragment to be integrated stably through generations in the cells of said plants, and wherein said protein has an enzymatic activity capable of causing inactivation or neutralization of said glutamine synthetase inhibitor.

A preferred DNA fragment is one derived from an antibiotic-producing-Streptomyces strain (or a sequence comprising a nucleotide sequence encoding the same activity) and which encodes resistance to —said GS

inhibitors.

Preferred nucleotide sequences for use in this invention encode a protein which has acetyl transferase activity with respect to said GS inhibitors.

A most preferred DNA fragment according to the invention comprises a nucleotide sequence coding for a polypeptide having a PPT acetyl transferase activity.

A particular DNA fragment according to the invention, for the subsequent transformation of plant cells, consists of a nucleotide sequence coding for at least part of a polypeptide having the following sequence :

X SER PRO GLU

183  
15 ARG ARG PRO ALA ASP ILE ARG ARG ALA THR GLU ALA ASP MET PRO

228  
ALA VAL CYS THR ILE VAL ASN HIS TYR ILE GLU THR SER THR VAL

273  
20 ASN PHE ARG THR GLU PRO GLN GLU PRO GLN GLU TRP THR ASP ASP

318  
LEU VAL ARG LEU ARG GLU ARG TYR PRO TRP LEU VAL ALA GLU VAL

363  
ASP GLY GLU VAL ALA GLY ILE ALA TYR ALA GLY PRO TRP LYS ALA

408  
25 ARG ASN ALA TYR ASP TRP THR ALA GLU SER THR VAL TYR VAL SER

453  
PRO ARG HIS GLN ARG THR GLY LEU GLY SER THR LEU TYR THR HIS

498  
30 LEU LEU LYS SER LEU GLU ALA GLN GLY PHE LYS SER VAL VAL ALA

543  
VAL ILE GLY LEU PRO ASN ASP PRO SER VAL ARG MET HIS GLU ALA

588  
LEU GLY TYR ALA PRO ARG GLY MET LEU ARG ALA ALA GLY PHE LYS

633  
35 HIS GLY ASN TRP HIS ASP VAL GLY PHE TRP GLN LEU ASP PHE SER

678  
LEU PRO VAL PRO PRO ARG PRO VAL LEU PRO VAL THR GLU ILE

723

in which X represents MET or VAL, which part of said polypeptide is of sufficient length to confer protection against Bialaphos to plant cells, when incorporated genetically and expressed therein, i.e. as termed  
 5 hereafter "plant-protecting capability" against Bialaphos.

A preferred DNA fragment consists of the following nucleotide sequence :

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10 183 184 185 186 187 188 189 190 191 192 193 194 195 196 197 198 199 200 201 202 203 204 205 206 207 208 209 210 211 212 213 214 215 216 217 218 219 220 221 222 223 224 225 226 227 228 229 230 231 232 233 234 235 236 237 238 239 240 241 242 243 244 245 246 247 248 249 250 251 252 253 254 255 256 257 258 259 260 261 262 263 264 265 266 267 268 269 270 271 272 273 274 275 276 277 278 279 280 281 282 283 284 285 286 287 288 289 290 291 292 293 294 295 296 297 298 299 300 301 302 303 304 305 306 307 308 309 310 311 312 313 314 315 316 317 318 319 320 321 322 323 324 325 326 327 328 329 330 331 332 333 334 335 336 337 338 339 340 341 342 343 344 345 346 347 348 349 350 351 352 353 354 355 356 357 358 359 360 361 362 363 364 365 366 367 368 369 370 371 372 373 374 375 376 377 378 379 380 381 382 383 384 385 386 387 388 389 390 391 392 393 394 395 396 397 398 399 400 401 402 403 404 405 406 407 408 409 410 411 412 413 414 415 416 417 418 419 420 421 422 423 424 425 426 427 428 429 430 431 432 433 434 435 436 437 438 439 440 441 442 443 444 445 446 447 448 449 450 451 452 453 454 455 456 457 458 459 460 461 462 463 464 465 466 467 468 469 470 471 472 473 474 475 476 477 478 479 480 481 482 483 484 485 486 487 488 489 490 491 492 493 494 495 496 497 498 499 500 501 502 503 504 505 506 507 508 509 510 511 512 513 514 515 516 517 518 519 520 521 522 523 524 525 526 527 528 529 530 531 532 533 534 535 536 537 538 539 540 541 542 543 544 545 546 547 548 549 550 551 552 553 554 555 556 557 558 559 560 561 562 563 564 565 566 567 568 569 570 571 572 573 574 575 576 577 578 579 580 581 582 583 584 585 586 587 588 589 590 591 592 593 594 595 596 597 598 599 600 601 602 603 604 605 606 607 608 609 610 611 612 613 614 615 616 617 618 619 620 621 622 623 624 625 626 627 628 629 630 631 632 633 634 635 636 637 638 639 640 641 642 643 644 645 646 647 648 649 650 651 652 653 654 655 656 657 658 659 660 661 662 663 664 665 666 667 668 669 670 671 672 673 674 675 676 677 678 679 680 681 682 683 684 685 686 687 688 689 690 691 692 693 694 695 696 697 698 699 700 701 702 703 704 705 706 707 708 709 710 711 712 713 714 715 716 717 718 719 720 721 722 723 724 725 726 727 728 729 730 731 732 733 734 735 736 737 738 739 740 741 742 743 744 745 746 747 748 749 750 751 752 753 754 755 756 757 758 759 760 761 762 763 764 765 766 767 768 769 770 771 772 773 774 775 776 777 778 779 780 781 782 783 784 785 786 787 788 789 790 791 792 793 794 795 796 797 798 799 800 801 802 803 804 805 806 807 808 809 810 811 812 813 814 815 816 817 818 819 820 821 822 823 824 825 826 827 828 829 830 831 832 833 834 835 836 837 838 839 840 841 842 843 844 845 846 847 848 849 850 851 852 853 854 855 856 857 858 859 860 861 862 863 864 865 866 867 868 869 870 871 872 873 874 875 876 877 878 879 880 881 882 883 884 885 886 887 888 889 890 891 892 893 894 895 896 897 898 899 900 901 902 903 904 905 906 907 908 909 910 911 912 913 914 915 916 917 918 919 920 921 922 923 924 925 926 927 928 929 930 931 932 933 934 935 936 937 938 939 940 941 942 943 944 945 946 947 948 949 950 951 952 953 954 955 956 957 958 959 960 961 962 963 964 965 966 967 968 969 970 971 972 973 974 975 976 977 978 979 980 981 982 983 984 985 986 987 988 989 990 991 992 993 994 995 996 997 998 999 1000

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35 or of a part thereof expressing a polypeptide having

plant-protecting capability against Bialaphos.

The invention also relates to any DNA fragment differing from the preferred one indicated hereabove by the replacement of any of its nucleotides by others, yet without modifying the genetic information of the preferred DNA sequence mentioned hereabove (normally within the meaning of the universal genetic code), and furthermore to any equivalent DNA sequence which would encode a polypeptide having the same properties, particularly a Bialaphos-resistance-activity.

It will be understood that the man skilled in the art should be capable of readily assessing those parts of the nucleotide sequences that could be removed from either side of any of the DNA fragments according to the invention, for instance by removing terminal parts on either side of said DNA fragment, such as by an exonucleolytic enzyme, for instance Bal31, by recloning the remaining fragment in a suitable plasmid and by assaying the capacity of the modified plasmid to transform appropriate cells and to protect it against the Bialaphos antibiotic or herbicide as disclosed later, whichever assay is appropriate.

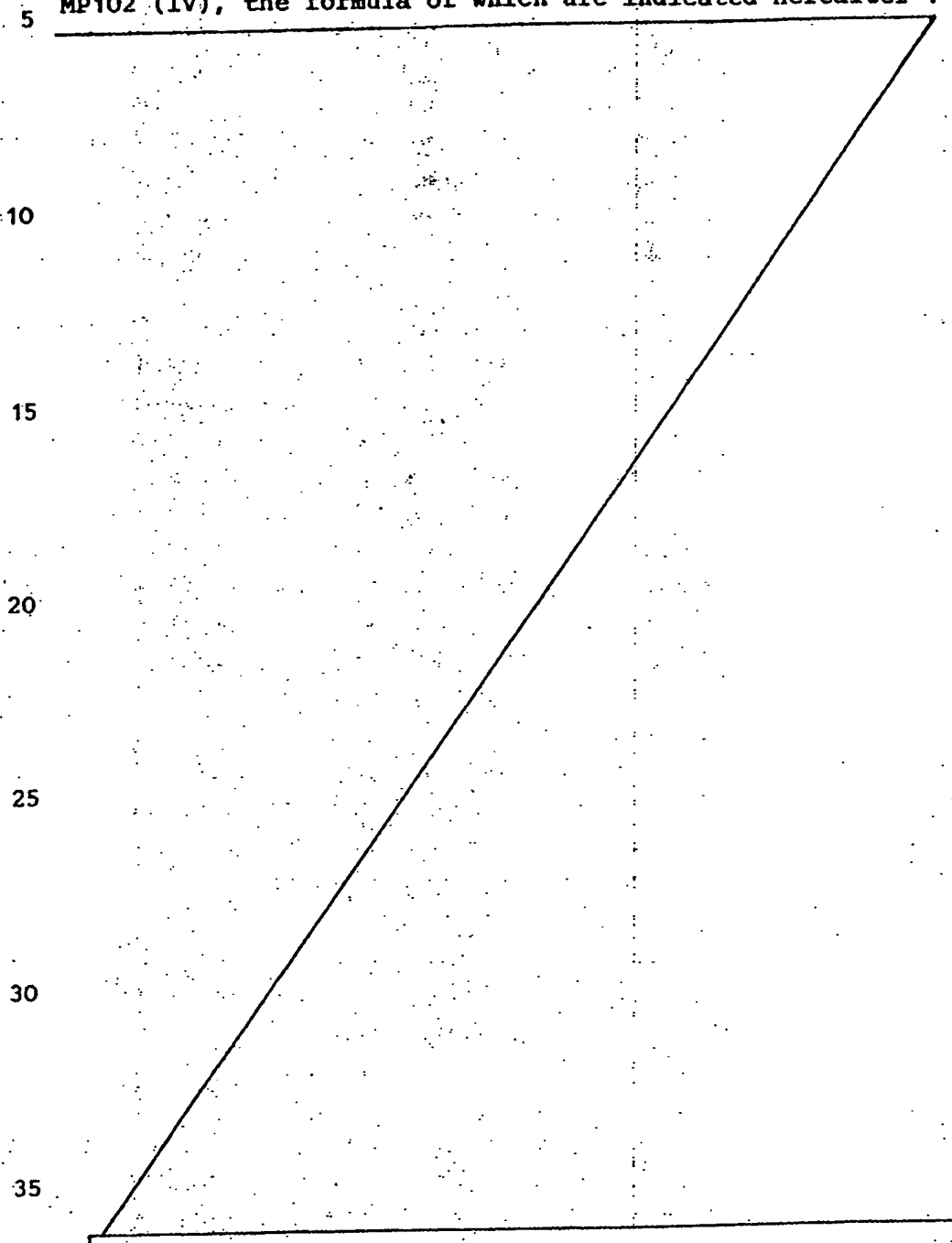
For the easiness of language, these DNA fragments will be termed hereafter as "Bialaphos-resistance DNA". In a similar manner, the corresponding polypeptide will be termed as "Bialaphos-resistance enzyme".

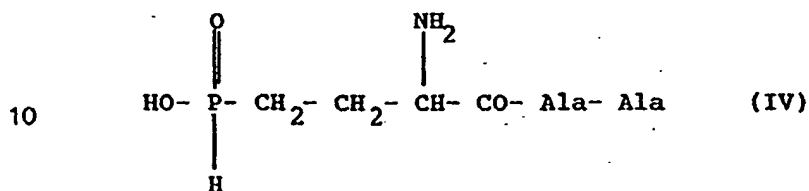
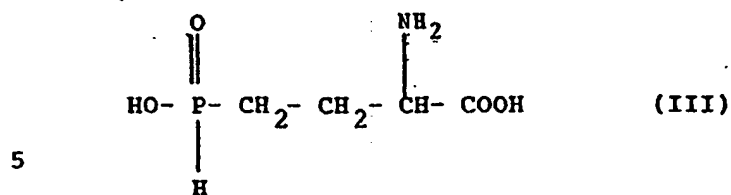
While in the preceding discussion particular emphasis has been put on DNA fragments capable, when introduced into plant cells and plants, to confer on them protection against Bialaphos or PPT, it should be understood that the invention should in no way be deemed as limited thereto.

In a same manner, the invention pertains to DNA fragments which, when introduced into such plant cells, would also confer on them a protection against other GS



inhibitors, for instance of intermediate products involved in the natural biosynthesis of phosphinotricin, such as the compounds designated by the abbreviations MP101 (III), MP102 (IV), the formula of which are indicated hereafter :





15 More generally, the invention has opened the route to the production of DNA fragments which, upon proper incorporation into plant cells and plants, can protect them against GS inhibitors when contacted therewith, as this will be shown in a detailed manner in relation to Bialaphos and PPT in the examples which will follow.

20 This having been established, it will be appreciated that any fragment encoding an enzymatic activity which would protect plant cells and plants against said GS inhibitors should be viewed as an equivalent of the preferred fragments which have been disclosed hereabove. This would apply especially to any  
25 DNA fragments that would result from genetic screening of the genomic DNAs of strains, particularly of antibiotic-producing strains, likely to possess genes which, even-though structurally different, would encode similar  
30 activity with respect to Bialaphos or PPT, or even with respect to other GS inhibitors. One might envisage similar genes in other strains producing a PPT derivative.

Therefore, it should be understood that the language "Bialaphos-resistance DNA" or "Bialaphos-resistance enzyme" used thereafter as a matter of  
35 convenience is intended to relate not only to the DNAs and

enzymes specifically concerned with resistance to PPT or most directly related derivatives, but more generally with other DNAs and enzymes which would be capable, under the same circumstances, of controlling the action in plants of GS inhibitors.

The invention also relates to DNA recombinants containing the above defined Bialaphos-resistance DNA fragments recombined with heterologous DNA, said heterologous DNA containing regulation elements and said Bialaphos-resistance DNA being under the control of said regulation elements in such manner as to be expressible in a foreign cellular environment compatible with said regulation elements.

By "heterologous DNA" is meant a DNA of an other origin than that from which said Bialaphos-resistance-DNA originated, e.g. is different from that of a Streptomyces hygroscopicus or Streptomyces viridochromogenes or even more preferably a DNA foreign to Streptomyces DNA. Particularly said regulation elements are those which are capable of controlling the transcription and translation of DNA sequences normally associated with them in said foreign environment. "Cellular" refers both to micro-organisms and to cell cultures.

This heterologous DNA may be a bacterial DNA, particularly when it is desired to produce a large amount of the recombinant DNA, such as for amplification purposes. In that respect a preferred heterologous DNA consists of DNA of E. coli or of DNA compatible with E. coli. It may be DNA of the same origin as that of the cells concerned or other DNA, for instance viral or plasmidic DNA known as capable of replicating in the cells concerned.

Preferred recombinant DNA contains heterologous DNA compatible with plant cells, particularly Ti-plasmid DNA.

Particularly preferred recombinants are those

which contain GS inhibitor inactivating DNA under the control of a promoter recognized by plant cells, particularly those plant cells on which inactivation of GS inhibitors is to be conferred.

Preferred recombinants according to the invention further relate to modified vectors, particularly plasmids, containing said GS-inhibitor-inactivating DNA so positioned with respect to regulation elements, including particularly promoter elements, that they enable said GS inhibitor-inactivating DNA to be transcribed and translated in the cellular environment which is compatible with said heterologous DNA. Advantageous vectors are those so engineered as to cause stable incorporation of said GS inhibitor-inactivating DNA in foreign cells, particularly in their genomic DNA. Preferred modified vectors are those which enable the stable transformation of plant cells and which confer to the corresponding cells, the capability of inactivating GS inhibitors.

It seems that, as described later, the initiation codon of the Bialaphos-resistance-gene of the Streptomyces hygroscopicus strain used herein is a GTG codon. But in preferred recombinant DNAs or vectors, the Bialaphos-resistance-gene is modified by substitution of an ATG initiation codon for the initiation codon GTG, which ATG enables translation initiation in plant cells.

In the example which follows, the plant promoter sequence which has been used was constituted by a promoter of the 35 S cauliflower mosaic virus. Needless to say that the man skilled in the art will be capable of selecting other plant promoters, when more appropriate in relation to the plant species concerned.

According to an other preferred embodiment of the invention, particularly when it is desired to achieve transport of the enzyme encoded by the Bialaphos-resistance-DNA into the chloroplasts, the heterologous DNA

fragment is fused to a gene or DNA fragment encoding a transit peptide, said last mentioned fragment being then intercalated between the GS inhibitor inactivating gene and the plant promoter selected.

As concerns means capable of achieving such constructions, reference can be made to the following British applications 84 32757 filed on December 28, 1984 and 85 00336 filed on January 7, 1985 and to the related applications filed in the United-States of America (n° 755,173, filed on July 15, 1985), in the European Patent Office (n° 85 402596.2, filed on December 20, 1985), in Japan (n° 299 730, filed on December 27, 1985), in Israel (n° 77 466 filed on December 27, 1985) and in Australia (n° 5 165 485, filed on December 24, 1985), all of which are incorporated herein by reference.

Reference can also be made to the scientific literature, particularly to the following articles :

- VAN DEN BROECK et al., 1985, Nature, 313, 358-363 ;
- SCHREIER and al., Embo. J., vol. 4, n° 1, 25-32.

These articles are also incorporated herein by reference.

For the sake of the record, be it recalled here that under the expression "transit peptide", one refers to a polypeptide fragment which is normally associated with a chloroplast protein or a chloroplast protein sub-unit in a precursor protein encoded by plant cell nuclear DNA. The transit peptide then separates from the chloroplast protein or is proteolitically removed, during the translocation process of the latter protein into the chloroplasts. Examples of suitable transit peptides are those associated with the small subunit of ribulose-1,5 biphosphate (RuBP) carboxylase or that associated with the chlorophyl a/b binding proteins.

There is thus provided DNA fragments and DNA

recombinants which are suitable for use in the process defined hereafter.

More particularly the invention also relates to a process, which can be generally defined as a process for producing plants and reproduction material of said plants including a heterologous genetic material stably integrated therein and capable of being expressed in said plants or reproduction material in the form of a protein capable of inactivating or neutralizing the activity of a glutamine synthetase-inhibitor, comprising the non biological steps of producing plants cells or plant tissue including said heterologous genetic material from starting plant cells or plant tissue not able to express that inhibiting or neutralizing activity, regenerating plants or reproduction material of said plants or both from said plant cells or plant tissue including said genetic material and, optionally, biologically replicating said last mentioned plants or reproduction material or both, wherein said non-biological steps of producing said plant cells or plant tissue including said heterologous genetic material, comprises transforming said starting plant cells or plant tissue with a DNA-recombinant containing a nucleotide sequence encoding said protein, as well as the regulatory elements selected among those which are capable of enabling the expression of said nucleotide sequence in said plant cells or plant tissue, and to cause the stable integration of said nucleotide sequence in said plant cells and tissue, as well as in the plant and reproduction material processed therefrom throughout generations.

The invention also relates to the cell cultures containing Bialaphos-resistance-DNA, or more generally said GS-inhibitor-inactivating DNA, which cell cultures have the property of being resistant to a composition containing a GS inhibitor, when cultured in a medium containing a such composition at dosages which would be

destructive for non transformed cells.

The invention concerns more particularly those plant cells or cell cultures in which the Bialaphos-resistance DNA is stably integrated and which remains present over successive generations of said plant cells. Thus the resistance to a GS inhibitor, more particularly Bialaphos or PPT, can also be considered as a way of characterizing the plant cells of this invention.

Optionally one may also resort to hybridization experiments between the genomic DNA obtained from said plant cells with a probe containing a GS inhibitor inactivating DNA sequence.

More generally the invention relates to plant cells, reproduction material, particularly seeds, as well as plants containing a foreign or heterologous DNA fragment stably integrated in their respective genomic DNAs, said fragments being transferred throughout generations of such plant cells, reproduction material, seeds and plants, wherein said DNA fragment encodes a protein inducing a non-variety-specific enzymatic activity capable of inactivating or neutralizing GS inhibitors, particularly Bialaphos and PPT, more particularly to confer on said plant cells, reproduction material, seeds and plants a corresponding non-variety-specific phenotype of resistance to GS inhibitors.

"Non-variety-specific" enzymatic activity or phenotype aims at referring to the fact that they are not characteristic of specific plant genes or species as this will be illustrated in a non-limitative way by the examples which will follow. They are induced in said plant materials by essentially non-biological processes applicable to plants belonging to species normally unrelated with one another and comprising the incorporation into said plant material of heterologous DNA, e.g. bacterial DNA or chemically synthesized DNA, which does

not normally occur in said plant material or which normally cannot be incorporated therein by natural breeding processes, and which yet confers a common phenotype (e.g. herbicide resistance) to them.

The invention is of particular advantageous use in processes for protecting field-cultivated plant species against weeds, which processes comprise the step of treating the field with an herbicide, e.g. Bialaphos or PPT in a dosage effective to kill said weeds, wherein the cultivated plant species then contain in their genome a DNA fragment encoding a protein having an enzymatic activity capable of neutralizing or inactivating said GS inhibitor.

By way of illustration only, effective doses for use in the abovesaid process range from about 0.4 to about 1.6 kg/Hectare of Bialaphos or PPT.

There follows now a disclosure of how the preferred DNA fragment described hereabove was isolated starting from the Streptomyces hygrosopicus strain available at the American Type Culture Collection under deposition number ATCC 21 705, by way of exemplification only.

The following disclosure also provides the technique which can be applied to other strains producing compounds with a PPT moiety.

The disclosure will then be completed with the description of the insertion of a preferred DNA fragment conferring to the transformed cells the capability of inactivating Bialaphos and PPT. Thus the Bialaphos-inactivating-DNA fragment designated thereafter by Bialaphos-resistance gene or "sfr" gene, isolated by the above described technique into plasmids which can be used for transforming plant cells and conferring to them a resistance against Bialaphos, also merely by way of example for non-limitative illustration purposes.



The following disclosure is made with reference to the drawings in which :

5 - fig. 1 is a restriction map of a plasmid containing a Streptomyces hygrosopicus DNA fragment encoding Bialaphos-resistance, which plasmid, designated hereafter as pBG1 has been constructed according to the disclosure which follows ;

10 - fig. 2 shows the nucleotide sequence of a smaller fragment obtained from pBG1, subcloned into another plasmid (pBG39) and containing the resistance gene ;

15 - fig. 3 shows the construction of a series of plasmids given by way of example, which plasmids aim at providing suitable adaptation means for the insertion therein of the Bialaphos-resistance gene or "sfr" gene ;

20 - fig. 4A and 4B show the construction of a series of plasmids given by way of example, which plasmids contain suitable plant cell promoter sequences able to initiate transcription and expression of the foreign gene inserted under their control into said plasmids ;

- fig. 5A shows a determined fragment of the nucleotide sequence of the plasmid obtained in figure 3 ;

25 - fig. 5B shows the reconstruction of the first codons of a Bialaphos-resistance gene, from a FokI/BglII fragment obtained from pBG39 and the substitution of an ATG initiation codon for the GTG initiation codon of the natural "sfr" gene ;

30 - fig. 5C shows the reconstruction of the entire "sfr" gene, namely the last codons thereof, and its insertion into a plasmid obtained in figures 4A and 4B ;

- fig. 6A shows an expression vector containing the "sfr" gene placed under the control of a plant cell promoter ;

35 - fig. 6B shows another expression vector deriving from the one shown in fig. 6A, by the substitution of some nucleotides.

- fig. 7 shows the construction of a series of plasmids given by way of examples, to ultimately produce plasmids containing the promoter region and the transit peptide sequence of a determined plant cell gene, for the insertion of the "sfr" gene under the control of said promoter region and downstream of said transit peptide sequence.

The following experiment was set up to isolate a Bialaphos-resistance-gene from S. hygrosopicus, according to standard techniques for cloning into Streptomyces.

2.5  $\mu\text{g}$  of S. hygrosopicus genomic DNA and 0.5  $\mu\text{g}$  of Streptomyces vector pIJ61 were cleaved with PstI according to the method described in ref. 6. The vector fragments and genomic fragments were mixed and ligated (4 hours at 10°C followed by 72 hours at 4°C in ligation salts which contain 66 mM Tris-HCl (pH 7.5), 1 mM EDTA, 10 mM  $\text{MgCl}_2$ , 10 mM 2-mercaptoethanol and 0.1 mM ATP) at a total DNA concentration of 40  $\mu\text{g ml}^{-1}$  with T4 DNA ligase. Ligation products were introduced into  $3 \times 10^9$  S. lividans strain 66 protoplasts by a transformation procedure mediated by polyethylene-glycol (PEG) as described hereafter. These protoplasts gave rise to  $5 \times 10^7$  colonies and  $4 \times 10^4$  pocks after regeneration on 20 plates of R2 agar containing 0.5 % of Difco yeast extract (R2 YE). Preparation and composition of the different mediums and buffers used in the disclosed experiments are described hereinafter. When these lawns were replica-plated on minimal medium plates containing 50  $\mu\text{g ml}^{-1}$  Bialaphos, drug resistant colonies appeared at a frequency of 1 per  $10^4$  transformants. After purification of the drug resistant colonies, there plasmid DNA was isolated and used to retransform S. lividans protoplasts. Non selective regeneration followed by replication to Bialaphos-containing-medium demonstrated a 100 % correlation between pocks and Bialaphos resistant growth. The recombinant plasmids of

several resistant clones all contained a 1.7 Kb PstI insert (see fig. 1).

Subcloning of the herbicide resistance gene

5 The 1.7 Kb PstI insert was then subcloned into the high copy number streptomycete vector pIJ385 to generate plasmid pBG20. S. lividans strains which contained pBG20 were more than 500 times more resistant to Bialaphos.

10 S. lividans growth is normally inhibited in minimal medium containing 1 µg/ml Bialaphos; growth of transformants containing pBG20 was not noticeably inhibited in a medium containing 500 µg/ml Bialaphos. The PstI fragment was also subcloned in either orientation into the PstI site of the plasmid pBR322, to produce plasmids pBG1 and pBG2, according to their orientation. A test on minimal M9 medium  
15 demonstrated that E. coli E8767 containing pBG1 or pBG2 was resistant to Bialaphos.

A 1.65 Kb PstI - BamHI fragment was subcloned from pBG1 into the plasmid pUC19 to produce the plasmid  
20 pBG39, and conferred Bialaphos resistance to E. coli, W3110, C600 and JM83.

Using an in vitro coupled transcription-translation system (ref. 5) from S. lividans extracts, the  
25 1.65 Kb PstI - BamHI fragment in pBG39 was shown to direct the synthesis of a 22 Kd protein. In the following, this 1.65 Kb insert includes a fragment coding for a 22 Kd protein and will be called "sfr" gene.

Fine mapping and sequencing of the gene

30 A 625 bp Sau3A fragment was subcloned from pBG39 into pUC19 and still conferred Bialaphos resistance to a E. coli W3110 host. The resulting clones were pBG93 and pBG94, according to the orientation.

The orientation of the gene in the Sau3A fragment was indicated by experiments which have shown that  
35 Bialaphos resistance could be induced with IPTG from the pUC19 lac promoter in pBG93. In the presence of IPTG

(0.5 mM) the resistance of pBG93/W3110 increased from 5 to 50 µg/ml on a M9 medium containing Bialaphos. The W3110 host devoid of pBG93, did not grow on M9 medium containing 5 µg/ml Bialaphos. These experiments demonstrated that the Sau3A fragment could be subcloned without loss of activity. They also provided for the proper orientation as shown in the fig. 2, enclosed thereafter. The protein encoded by these clones was detected by using coupled transcription-translation systems derived from extracts of S. lividans (ref. 7). Depending on the orientation of the Sau3A fragment, translation products of different sizes were observed ; 22 Kd for pBG94 and ± 28 Kd for pBG93. This indicated that the Sau3A fragment did not contain the entire resistance gene and that a fusion protein was formed which included a polypeptide sequence resulting from the translation of a pUC19 sequence.

In order to obtain large amounts of the protein, a 1.7 Kb PstI fragment from pBG1 was cloned into the high copy number Streptomyces replicon pIJ385. The obtained plasmid, pBG20, was used to transform S. hygroscopicus. Transformants which contained this plasmid had more than 5 times as much PPT acetylating activity and also had increased amounts of a 22 kd protein on sodium dodecyl-sulfate gels (SDS gels). Furthermore, both the acetyl transferase and the 22 kd protein appeared when the production of Bialaphos begun. The correlation of the in vitro data, kinetics of synthesis, and amplified expression associated with pBG20 transformants strongly implied that this 22 Kd band was the gene product.

The complete nucleotide sequence of the 625 bp Sau3A fragment was determined as well as of flanking sequences. Computer analysis revealed the presence of an open reading frame over the entire length of the Sau3A fragment.

#### Characterization of the *sfr* gene product

A series of experiments were performed to determine that the open reading frame of the "sfr" gene indeed encoded the Bialaphos resistance enzyme. To determine the 5' end of the resistance gene, the NH<sub>2</sub>-terminal sequence of the enzyme was determined. As concerns more particularly the technique used to determine the said sequence, reference is made to the technique developed by J. VANDEKERCKHOVE, Eur. J. Bioc. 152, p. 9-19, 1985, and to French patent applications n° 85 14579 filed on October 1st, 1985 and n° 85 13046 filed on September 2nd, 1985, all of which are incorporated herein by reference.

This technique allows the immobilization on glass fibre sheets coated with the polyquaternary amine commercially available under the registered trademark POLYBRENE of proteins and of nucleic acids previously separated on a sodium dodecylsulfate containing polyacrylamide gel. The transfer is carried out essentially as for the protein blotting on nitrocellulose membranes (ref. 8). This allows the determination of amino-acid composition and partial sequence of the immobilized proteins. The portion of the sheet carrying the immobilized 22 kd protein produced by *S. hygrosopicus* pBG20 was cut out and the disc was mounted in the reaction chambre of a gas-phase sequenator to subject the glass-fibre bound 22 Kd protein to the Edman degradation procedure. The following amino-acid sequence was obtained :

Pro-Glu-Arg-Arg-Pro-Ala-Asp-Ile-Arg-Arg

This sequence matched an amino-acid sequence which was deduced from the open reading frame of the 625 bp *Sau*3A fragment. It corresponded to the stretch from codon 3 to codon 12.

Thus, the NH<sub>2</sub>-terminus of the 22 Kd protein was upstream of this sequence. It was determined that translation of the actual protein was likely to be initiated 2 amino-acids earlier at a GTG initiation codon. GTG is

often used as initiator codon in *Streptomyces* and translated as methionine. The protein translated from the GTG initiation codon would be 183 amino-acids long and would  
5 have a molecular weight of 20 550. This was in good agreement with the observed approximate molecular weight of 22 000.

Furthermore, the termination codon, TGA, was located just downstream of the Sau3A site. Cloning of the  
10 625 bp Sau3A fragment in a BamHI site digested pUC19 did not result in the reconstruction of the termination codon. This explained the fusion proteins which were observed in the in vitro transcription-translation analysis.

#### Mechanism of PPT-resistance

15 Having defined a first phenotype and some of the physical characteristics of the resistance gene and its gene product, a series of experiments was then carried out to understand the mechanism by which it confers resistance. As described hereabove, PPT is the portion of  
20 Bialaphos which inhibits glutamine synthetase (GS) and that N-acetyl PPT is not an inhibitor. Using a standard assay (ref. 9), S. hygrosopicus ATCC 21 705 derivatives were shown to contain a PPT acetyl transferase which was not found in S. lividans. The activity does not acetylate  
25 the Bialaphos tripeptide. S. lividans carrying the resistance gene cloned in pBG20 or pBG16 (a plasmid containing the 625 bp Sau3A fragment cloned into another streptomycete vector, pIJ680) also contained the activity which could acetylate PPT but not Bialaphos. The PPT derived reaction product produced by extracts of pBG20/  
30 S. lividans was isolated in order to confirm that it was indeed acetyl-PPT. Analysis by mass spectroscopy showed that the molecular weight had increased relative to PPT by the equivalent of one acetyl group. It was thus concluded  
35 that the 625 bp Sau3A fragment contained sequences which code for PPT acetyl transferase.

The experimental conditions and reagents used in the techniques disclosed hereabove were as follows :

Preparation and composition of the mediums and buffers above used

5 1° P medium : 10.3 g of sucrose, 0.025 g of  $K_2SO_4$ , 0.203 g of  $MgCl_2 \cdot 6H_2O$  and 0.2 ml of a trace element solution are dissolved in 80 ml of distilled water and autoclaved. Then in order, 1 ml of  $KH_2PO_4$  (0.5 %), 10 ml of  $CaCl_2 \cdot 2H_2O$  (3.68 %), and 10 ml of TES buffer (0.25 M),  
10 pH : 7.2) are added. Trace element solution (per litre) :  $ZnCl_2$ , 40 mg ;  $FeCl_3 \cdot 6H_2O$ , 200 mg ;  $CuCl_2 \cdot 2H_2O$ , 10 mg ;  $MnCl_2 \cdot 4H_2O$ , 10 mg ;  $Na_2B_4O_7 \cdot 10H_2O$ , 10 mg ;  $(NH_4)_6Mo_7O_{24} \cdot 4H_2O$ , 10 mg.

15 2° R2YE : 10.3 g of sucrose, 0.025 g of  $K_2SO_4$ , 1.012 g of  $MgCl_2 \cdot 6H_2O$ , 1 g of glucose, 0.01 g of Difco casamino acids, and 2.2 g of Difco agar are dissolved in 80 ml distilled water and autoclaved. 0.2 ml of trace element solution, 1 ml of  $KH_2PO_4$  (0.5 %), 8.02 ml of  $CaCl_2 \cdot 2H_2O$  (3.68 %), 1.5 ml of L-proline (20 %), 10 ml of  
20 TES buffer (0.25 M) (pH : 7.2), 0.5 ml of (1 M) NaOH, 5 ml of yeast extract (10 %) are sequentially added.

3° TE : 10 mM TRIS HCl, 1 mM EDTA, pH 8.0.

4° YEME : Difco yeast extract (0.3 %), Difco peptone (0.5 %), oxoid malt extract (0.3 %), glucose (1 %).

Transformation of *S. lividans* protoplasts

1. A culture composed of 25 ml YEME, 34 % sucrose, 0.005 M  $MgCl_2$ , 0.5 % glycine, in a 250 ml baffled flask, is centrifuged during 30 to 36 hours.

30 2. The pellet is suspended in 10.3 % sucrose and centrifuged. This washing is repeated once.

3. The mycelium is suspended in 4 ml lysozyme solution (1 mg/ml in P medium with  $CaCl_2$  and  $MgCl_2$  concentrations reduced to 0.0025 M) and incubated at 30°C for 15 to  
35 60 minutes.

4. The solution is mixed by pipetting three times in a

5 ml pipette and incubated for further 15 minutes.

5. P medium (5 ml) is added and mixed by pipetting as in step 4.

5 6. The solution is filtered through cotton wool and protoplasts are gently sedimented in a bench centrifuge at 800 x G during 7 minutes.

7. Protoplasts are suspended in 4 ml P medium and centrifuged again.

10 8. Step 7 is repeated and protoplasts are suspended in the drop of P medium left after pouring off the supernatant (for transformation).

9. DNA is added in less than 20 µl TE.

10. 0.5 ml PEG 1 000 solution (2.5 g PEG dissolved in  
15 7.5 ml of 2.5 % sucrose, 0.0014 K<sub>2</sub>SO<sub>4</sub>, 0.1 M CaCl<sub>2</sub>, 0.05 M TRIS-maleic acid, pH 8.0, plus trace elements) is immediately added and pipetted once to mix the components.

11. After 60 seconds, 5 ml of P medium are added and the protoplasts are sedimented by gentle centrifugation.

20 12. The pellet is suspended in P medium (1 ml).

13. 0.1 ml is plated out on R2YE plates (for transformation dry plates to 85 % of their fresh weight e. g. in a laminar flow cabinet).

14. Incubation at 30°C.

25 A - Construction of a "sfr" gene cassette

A "sfr" gene cassette was constructed to allow subsequent cloning in plant expression vectors.

Isolation of a FokI-BglII fragment from the plasmid pBG39 containing a "sfr" gene fragment led to the loss  
30 of the first codons, including the initiation codon, and of the last codons, including the stop codon.

This fragment of the "sfr" gene could be reconstructed in vitro with synthetic oligonucleotides which encode appropriate amino-acids.

35 The complementary synthetic oligonucleotides were 5'-CATGAGCCCAGAAC and 3'-TCGGGTCTTGCTGC.



By using such synthetic oligonucleotides, the 5' end of the "sfr" gene could be reformed and the GTG initiation codon substituted for a codon well translated by plant cells, particularly an ATG codon.

The DNA fragment containing the oligonucleotides linked to the "sfr" gene was then inserted into an appropriate plasmid, which contained a determined nucleotide sequence thereafter designated by an "adapter" fragment.

10 This adapter fragment comprised :

- a TGA termination codon which enabled the last codons of the "sfr" gene to be reformed ;

- appropriate restriction sites which enabled the insertion of the fragment of the nucleotide sequence comprising the "sfr" gene partially reformed with the synthetic oligonucleotides ; this insertion resulted in the reconstruction of an intact "sfr" gene ;

- appropriate restriction sites for the isolation of the entire "sfr" gene.

20 The "sfr" gene was then inserted into another plasmid, which contained a suitable plant promoter sequence. The plant promoter sequence consisted of the cauliflower mosaic virus promoter sequence (p35S). Of course the invention is not limited to the use of this particular promoter. Other sequences could be chosen as promoters suitable in plants, for example the TR 1'-2' promoter region and the promoter fragment of a Rubisco small subunit gene from Arabidopsis thaliana hereafter described.

#### 30 1' Construction of the plasmid pLK56.2 (fig. 3)

The construction of plasmid pLK56.2 aimed at obtaining a suitable adaptor including the following sequence of restriction sites : SmaI, BamHI, NcoI, KpnI, BglII, MluI, BamHI, HindIII and XbaI.

35 The starting plasmids used for this construction, pLK56, pJB64 and pLK33 were those disclosed by BOTTERMAN

(ref. 11).

The DNA fragments hereafter described were isolated and separated from low melting point agarose (LGA).

5 The plasmid pLK56 was cleaved by the enzymes BamHI and NdeI. A NcoI-NdeI fragment (referred to in the drawings by arc "a" in broken line) obtained from plasmid pJB64 was substituted in pLK56 for the BamHI-NdeI fragment shown at "b". Ligation was possible after filling in the  
10 BamHI and NcoI protruding ends with the DNA polymerase I of E. coli (Klenow's fragment).

Particularly recircularization took place by means of a T4 DNA ligase. A new plasmid pLK56.3 was obtained.

This plasmid was cleaved by the enzymes XbaI and  
15 PstI.

The BamHI-PstI fragment of pLK33 (c) (on fig. 3) was substituted for the XbaI-PstI fragment (d) of pLK56.3, after repairing of their respective ends by Klenow's fragment.

20 After recircularization by means of the T4 DNA ligase, the obtained plasmid pLK56.2 contained a nucleotide sequence which comprised the necessary restriction sites for the subsequent insertion of the "sfr" gene.

2' Construction of the plasmid pGSH150 (fig. 4A)

25 Parallel with the last discussed construction, there was produced a plasmid containing a promoter sequence recognized by the polymerases of plant cells and including suitable restriction sites, downstream of said promoter sequence in the direction of transcription, which  
30 restriction sites are then intended to enable the accommodation of the "sfr" gene then obtainable from pLK56.2, under the control of said plant promoter.

Plasmid pGV825 is described in DEBLAERE et al. (ref. 10). Plasmid pJB63 is from BOTTERMAN (ref. 11).

35 pGV825 was linearized with PvuII and recircularized by the T4 DNA ligase, resulting in the deletion of

an internal PvuII fragment shown at (e), (plasmid pGV956).

pGV956 was then cleaved by BamHI and BglII.

5 The BamHI-BglII fragment (f) obtained from pJB63 was dephosphorylated with calf intestine phosphatase (CIP) and substituted for the BamHI-BglII fragment of pGV956.

Plasmid pGV1500 was obtained after recircularization by means of T4 DNA ligase.

10 An EcoRI-HindIII fragment obtained from plasmid pGSH50 was purified. The latter plasmid carried the dual TR 1'-2' promoter fragment described in VELTEN et al., (ref.13). This fragment was inserted in pGV1500, digested with HpaI and HindIII and yielded pGSH150.

15 This plasmid contains the promoter fragment in front of the 3' end of the T-DNA transcript 7 and a BamHI and ClaI sites for cloning.

3' Construction of the plasmid pGSJ260 (fig. 4B)

20 CP3 is a plasmid derived from pBR322 and which contains the 35S promoter region of cauliflower mosaic virus within a BamHI fragment.

pGSH150 was cut by BamHI and BglII.

25 The BamHI-BglII fragment (h) of CP3, which contained the nucleotide sequence of p35S promoter, was substituted for the BamHI-BglII fragment (i) in pGSH150 to form plasmid pGSJ250. pGSJ250 was then opened at its BglII restriction site.

A BamHI fragment obtained from mGV2 (ref. 12) was inserted in pGSJ250 at the BglII site to form plasmid pGSJ260.

30 However prior to inserting the "sfr" gene obtainable from pLK56.2 into plasmid pGSJ260, it was still desirable to further modify the first in order to permit insertion in a more practical manner. Thus pLK56.2 was further modified as discussed below to yield pGSR1.

35 Starting from plasmid pGSJ260, two plasmid constructions for subsequent transformations of plant cells

were made :

- a first plasmid permitting the expression of the "sfr" gene in the cytoplasm of plant cells, and

5       - a second plasmid so modified as to achieve transport of the Bialaphos-resistance enzymes to the chloroplasts of plant cells.

First case : plasmid enabling the expression of the "sfr" gene in the cytoplasm of plant cells

10 Cloning of the sfr gene cassette in a plant expression vector (pGSR2) (fig. 5)

On figure 5A, the nucleotide sequence of the adapter of pLK56.2 is shown. In particular, the locations of BamHI, NcoI, BglII restriction sites are shown.

15       This adapter fragment was cleaved by the enzymes NcoI and BglII.

Figure 5B shows the FokI-BglII fragment (j) obtained from pBG39. The locations of these two restriction sites are shown on figure 2.

20       Using synthetic oligonucleotides, the first codons of the "sfr" gene were reformed, particularly the 5' end of the gene in which a ATG initiation codon was substituted for the initial GTG codon.

25       This FokI-BglII fragment completed with the synthetic oligonucleotides was then substituted in pLK56.2 for the NcoI-BglII fragment of the adapter. The 3' end of the gene was thus reformed too, after recircularization with T4 DNA ligased. The plasmid obtained, pGSR1, thus contained the entire "sfr" gene inserted in its adapter.

30       The plasmid pGSJ260 was then opened by BamHI (fig. 5C) and the BamHI fragment obtained from pGSR1, which contained the entire "sfr" gene, was inserted into pGSJ260.

35       The obtained plasmid, pGSR2 (see fig. 6A) contained a pBR322 replicon, a bacterial streptomycin resistance gene (SDM-SP-AD-transferase) and an engineered

T-DNA consisting of :

- the border fragments of the T-DNA ;
- a chimeric kanamycin gene which provided a dominant selectable marker in plant cells ; and

- a chimeric "sfr" gene.

The chimeric "sfr" gene consisting of :

- the promoter region of the cauliflower mosaic virus (p35S) ;
- the "sfr" gene cassette as described in fig. 5 ;
- the 3' untranslated region, including the polyadenylation signal of T-DNA transcript 7.

pGSR2 was introduced into Agrobacterium tumefaciens recipient C58ClRif<sup>R</sup> (pGV2260) according to the procedure described by DEBLAERE et al. (ref. 10).

This strain was used to introduce the chimeric "sfr" gene in N. tabacum SR<sub>1</sub> plants.

Two variant plasmids deriving from pGSR2, namely pGSFR280 and pGSFR281, have been constructed. They differ in the untranslated sequence following the transcription initiation site. In pGSR2, this fragment consists of the following sequence :

GAGGACACGCTGAAATCACCAGTCTCGGATCCATG ;

while it consists of :

GAGGACACGCTGAAATCACCAGTCTCTCTACAAATCGATCCATG

in pGSR280 and of

GAGGACACGCTGAAATCACCAGTCTCTCTACAAATCGATG

in pGSFR281, with an ATG codon being the initiation codon of the "sfr" gene. The "sfr" gene is also fused to the TR1'-2' promoter in the plasmid pGSH150 (fig. 4A) yielding pGSFR160 and pGSFR161 (fig. 6B). These plasmids contain slight differences in the pTR2 "sfr" gene configuration : the "sfr" gene is correctly fused to the endogenous gene 2' ATG in pGSFR161 (for sequences see ref. 13), whereas 4 extra base pairs (ATCC) are present just ahead of the ATG codon in pGSFR160. Otherwise, plasmids pGSFR161 and

PGSFR160 are completely identical.

All plasmids are introduced in Agrobacterium by cointegration in the acceptor plasmid pGV2260 yielding the  
5    respective plasmids PGSFR1280, PGSFR1281, PGSFR1160 and  
     PGSFR1161.

Second case : construction of a plasmid containing the  
     "sfr" gene downstream of a DNA sequence encoding a transit  
     peptide and suitable for achieving subsequent transloca-  
10    tion of the "sfr" gene expression product into plant-cell-  
     chloroplasts

In another set of experiments, the nucleotide  
sequence which contained the "sfr" gene was fused to a DNA  
sequence encoding a transit peptide so as to enable its  
15    transport into chloroplasts.

A fragment of the "sfr" gene was isolated from the  
adapter fragment above described and fused to a transit  
peptide. With synthetic oligonucleotides, the entire "sfr"  
gene was reconstructed and fused to a transit peptide.

20    The plasmid (plasmid PATS3 mentioned below) which  
     contained the nucleotide sequence encoding the transit  
     peptide comprised also the promoter sequence thereof.

Construction of the plasmid pGSR4 which contains the "sfr"  
     gene fused to a DNA sequence encoding transit peptide  
25    (fig. 7)

Plasmid pLK57 is from BOTTERMAN, (ref.11). Plasmid  
PATS3 is a pUC19 clone which contains a 2 Kb EcoRI genomic  
DNA fragment from Arabidopsis thaliana comprising the pro-  
moter region and the transit peptide nucleotide sequence  
30    of the gene, the expression thereof is the small subunit  
     of ribulose biphosphate carboxylase (ssu). The A. thaliana  
     small subunit was isolated as a 1 500 bp EcoRI-SphI frag-  
     ment. The SphI cleavage site exactly occurs at the site  
     where the coding region of the mature ssu protein starts.

35    Plasmids pLK57 and PATS3 were opened with EcoRI  
     and SphI. After recircularization by means of the T4 DNA

ligase, a recombinant plasmid pLKAB1 containing the sequence encoding the transit peptide (Tp) and its promoter region (Pssu) was obtained.

5 In order to correctly fuse the "sfr" gene at the cleavage site of the signal peptide, the N-terminal gene sequence was first modified. Since it was observed that N-terminal gene fusions with the "sfr" gene retain their enzymatic activity, the second codon (AGC) was modified to  
10 a GAC, yielding an NcoI site overlapping with the ATG initiator site. A new plasmid, pGSSFR2 was obtained. It only differs from pGSR1 (fig. 5B), by that mutation. The NcoI-BamHI fragment obtained from pGSR2 was fused at the SphI end of the transit peptide sequence. In parallel, the  
15 "sfr" gene fragment was fused correctly to the ATG initiator of the ssu gene (not shown in figures).

Introduction of the "sfr" gene into a different plant species

20 The Bialaphos-resistance induced in plants by the expression of chimeric genes, when the latter have been transformed with appropriate vectors containing said chimeric genes, has been demonstrated as follows. The recombinant plasmids containing the "sfr" gene were introduced separately by mobilization into Agrobacterium strain  
25 C58C<sub>1</sub> Rif<sup>R</sup> (pGV2260) according to the procedure described by DEBLAERE and al., Nucl. Acid. Res., 13, p. 1 477, 1985. Recombinant strains containing hybrid Ti plasmides were formed. These strains were used to infect and transform leaf discs of different plant species, according to a  
30 method essentially as described by HORSH and al., 1985, Science, vol. 227. Transformation procedure of these different plant species given by way of example, is described thereafter.

1. Leaf disc transformation of Nicotiana tabacum

Used Media are described thereafter :

5	A <sub>1</sub> MS salt/2	+ 1% sucrose 0.8 % agar pH 5.7
10	A <sub>10</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub> 750 mg/l CaCl <sub>2</sub> 2H <sub>2</sub> O 0.5 g/l 2-(N-Morpholino)ethane- sulfonic acid (MES) pH 5.7 30 g/l sucrose
15	A <sub>11</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub> 0.5 g/l MES pH 5.7 2 % glucose 0.8 % agar 40 mg/l adenine
20		+ 1 mg/l 6-Benzylaminopurine (BAP) 0.1 mg/l Indole-3-acetic acid (IAA) 500 mg/l Claforan
25	A <sub>12</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub> 0.5 g/l MES pH 5.7 2 % glucose 0.8 % agar 40 mg/l adenine
30		+ 1 mg/l BAP 200 mg/l claforan



32

A<sub>13</sub> MS-salt/2 + 3 % sucrose  
 0.5 MES g/l pH 5.7  
 0.7 % agar  
 200 mg/l claforan

5

Bacterial medium = min A : (Miller 1972) 60 mM

K<sub>2</sub>HPO<sub>4</sub> · 3H<sub>2</sub>O,

33 mM KH<sub>2</sub>PO<sub>4</sub> ; 7.5 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>

10

1.7 M trisodiumcitrat; 1 mM

MgSO<sub>4</sub> ;

2 g/l glucose ; 50 mg/l vita-  
 mine B1

15

- Plant material :

Nicotiana tabacum cv. Petit Havana SR1

Plants are used 6 to 8 weeks after subculture on  
 medium A<sub>1</sub>

- Infection :

20

- midribs and edges are removed from leaves.

- Remaining parts are cut into segments of about  
 0.25 cm<sup>2</sup> and are placed in the infection medium A<sub>10</sub> (about  
 12 segments in a 9 cm Petri dish containing 10 ml A<sub>10</sub>).

25

- Segments are then infected with 25 µl per Petri  
 dish of a late log culture of the Agrobacterium strain  
 grown in min A medium.

- Petri dish are incubated for 2 to 3 days at low  
 light intensity.

30

- After 2 to 3 days medium is removed and replaced  
 by 20 ml of medium A<sub>10</sub> containing 500 mg/l clarofan.

- Selection and shoot induction

- The leaf discs are placed on medium A<sub>11</sub> contain-  
 ing a selective agent :

100 mg/l kanamycin and

35

10 to 100 mg/l phosphinotricin.

- Leaf discs are transferred to fresh medium weekly.

5     - After 3 to 4 weeks regenerating calli arise. They are separated and placed on medium A<sub>12</sub> with the same concentration of selective agent as used for the selection.

- Rooting

10    - After 2 to 3 weeks the calli are covered with shoots, which can be isolated and transferred to rooting medium A<sub>13</sub> (without selection).

- Rooting takes 1 to 2 weeks.

- After a few more weeks, these plants are propagated on medium A<sub>1</sub>.

15    2. Tuber disc infection of Solanum tuberosum (potato)

Used media are described thereafter :

20     C<sub>1</sub> B5-medium     + 250 mg/l NH<sub>4</sub>NO<sub>3</sub>  
                           300 mg/l (CaH<sub>2</sub>PO<sub>4</sub>)<sub>2</sub>  
                           0.5 g/l MES pH 5.7  
                           0.5 g/l polyvinylpyrrolidone (PVP)  
                           40 g/l mannitol (=0.22M)  
 25                       0.8 % agar  
                           40 mg/l adenine

30     C<sub>2</sub> B5-medium     + 250 mg/l NH<sub>4</sub>NO<sub>3</sub>  
                           400 mg/l glutamine  
                           0.5 g/l MES pH 5.7  
                           0.5 g/l PVP  
                           40 g/l mannitol  
                           40 mg/l adenine  
 35                       0.8 % agar

		34	
		+ 0.5 mg/l transzeatine	
		0.1 mg/l IAA	
		500 mg/l clarofan	
5	C <sub>5</sub> MS salt/2	+ 3 % sucrose	
		0.7 % agar	
		pH 5.7	
10	C <sub>7</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub>	
		400 mg/l glutamine	
		0.5 g/l MES pH 5.7	
		0.5 g/l PVP	
		20 g/l mannitol	
15		20 g/l glucose	
		40 mg/l adenine	
		0.6 % agarose	
		+ 0.5 mg/l transzeatine	
20		0.1 mg/l IAA	
		500 mg/l clarofan	
	C <sub>8</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub>	
		400 mg/l glutamine	
25		0.5 g/l MES pH 5.7	
		0.5 g/l PVP	
		20 g/l mannitol	
		20 g/l glucose	
		40 mg/l adenine	
30		0.6 % agarose	
		+ 200 mg/l clarofan	
		1 mg/l transzeatine	

35

35

C<sub>9</sub> B5-medium + 250 mg/l NH<sub>4</sub>NO<sub>3</sub>  
400 mg/l glutamine  
0.5 g/l MES pH 5.7  
0.5 g/l PVP  
20 g/l mannitol  
20 g/l glucose  
40 mg/l adenine  
0.6 % agarose

5

10

+ 1 mg/l transzeatine  
0.01 mg/l Gibberellic acid A<sub>3</sub>  
(GA<sub>3</sub>)  
100 mg/l clarofan

15

C<sub>11</sub> MS salt/2 + 6 % sucrose  
0.7 % agar

Bacterial medium = min A : (Miller 1972 60 mM  $K_2HPO_4 \cdot 3H_2O$ ;  
33 mM  $KH_2PO_4$ ; 7.5 mM  $(NH_4)_2SO_4$ ;  
1.7 trisodiumcitrat; 1 mM  
 $MgSO_4$  ;  
2 g/l glucose; 50 mg/l vitami-  
ne B1.

25 - PLant material

Tubers of Solanum tuberosum c.v Berolina  
c.v Désirée

- Infection

- Potatoes are peeled and washed with water.

30           - Then they are washed with concentrated commercial bleach for 20 minutes, and

- rinsed 3 to 5 times with sterile water.

- The outer layer is removed (1 to 1.5 cm)

- The central part is cut into discs of about 1

35 cm<sup>2</sup> and 2 to 3 mm thick.

- Discs are placed on medium C<sub>1</sub> (4 pieces in a 9

cm Petri dish).

- 10  $\mu$ l of a late log culture of an Agrobacterium strain grown in min A medium is applied on each disc.

5       - Discs are incubated for 2 days at low light intensity.

- Selection and shoot induction

- Discs are dried on a filter paper and transferred to medium C<sub>2</sub> with 100 mg/l kanamycin.

10       - After one month small calli are removed from the discs and transferred to medium C<sub>7</sub> containing 50 mg/l kanamycin.

- After a few more weeks, the calli are transferred to medium C<sub>8</sub> containing 50 mg/l kanamycin.

15       - If little shoots start to develop, the calli are transferred to elongation medium C<sub>9</sub> containing 50 mg/l Kanamycin.

- Rooting

20       - Elongated shoots are separated and transferred to rooting medium C<sub>11</sub>.

- Rooted shoots are propagated on medium C<sub>5</sub>.

### 3. Leaf disc infection of Lycopersicum esculentum (tomato)

Used media are described thereafter

25       A<sub>1</sub> MS salt/2       + 1 % sucrose  
                                  0.8 % agar  
                                  pH 5.7

30       B<sub>1</sub> B5-medium       + 250 mg/l NH<sub>4</sub>NO<sub>3</sub>  
                                  0.5 g/l MES pH 5.7  
                                  0.5 g/l PVP  
                                  300 mg/l Ca (H<sub>2</sub>PO<sub>4</sub>)<sub>2</sub>  
                                  2 % glucose  
                                  40 mg/l adenine  
                                  40 g/l mannitol

35

		37
	B <sub>2</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub>
		0.5 g/l MES pH 5.7
		0.5 g/l PVP
5		400 mg/l glutamine
		2 % glucose
		0.6 % agarose
		40 mg/l adenine
		40 g/l mannitol
10		
		+ 0.5 mg/l transzeatine
		0.01 mg/l IAA
		500 mg/l claforan
15	B <sub>3</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub>
		0.5 g/l MES pH 5.7
		0.5 g/l PVP
		400 mg/l glutamine
		2 % glucose
20		0.6 % agarose
		40 mg/l adenine
		30 g/l mannitol
25		
		+ 0.5 mg/l transzeatine
		0.01 mg/l IAA
		500 mg/l clarofan
30	B <sub>4</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub>
		0.5 g/l MES pH 5.7
		0.5 g/l PVP
		400 mg/l glutamine
		2 % glucose
		0.6 % agarose
		40 mg/l adenine
35		20 g/l mannitol

		38	
		+ 0.5 mg/l transzeatine	
		0.01 mg/l IAA	
		500 mg/l clarofan	
5	B <sub>5</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub>	
		0.5 g/l MES pH 5.7	
		0.5 g/l PVP	
		400 mg/l glutamine	
10		2 % glucose	
		0.6 % agarose	
		40 mg/l adenine	
		10 g/l mannitol	
15		+ 0.5 mg/l transzeatine	
		0.01 mg/l IAA	
		500 mg/l clarofan	
20	B <sub>6</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub>	
		0.5 g/l MES pH 5.7	
		0.5 g/l PVP	
		400 mg/l glutamine	
		2 % glucose	
		0.6 % agarose	
25		40 mg/l adenine	
		+ 0.5 mg/l transzeatine	
		0.01 mg/l IAA	
		200 mg/l clarofan	
30	B <sub>7</sub> B5-medium	+ 250 mg/l NH <sub>4</sub> NO <sub>3</sub>	
		0.5 g/l MES pH 5.7	
		0.5 g/l PVP	
		400 mg/l glutamine	
35		2 % glucose	
		0.6 % agarose	

39

40 mg/l adenine

+ 1 mg/l transzeatine

5

200 mg/l clarofan

 $B_8$  MS salt/2

+ 2 % sucrose

0.5 g/l MES pH 5.7

0.7 % agar

10

 $B_9$  B5-medium+ 250 mg/l  $NH_4NO_3$ 

0.5 g/l MES pH 5.7

0.5 g/l PVP

2 % glucose

15

0.6 % agarose

40 mg/l adenine

+ 1 mg/l transzeatine

0.01 mg/l  $GA_3$ 

20

Bacterial medium = min A : (Miller 1972) 60 mM

 $K_2HPO_4 \cdot 3H_2O$  ;33 mM  $KH_2PO_4$ ; 7.5 mM  $(NH_4)_2SO_4$ ;

1.7 M trisodiumcitrat; 1 mM

25

 $MgSO_4$  ;

2 g/l glucose; 50 mg/l vitami-

ne B1

- Plant materialLycopersicum esculentum cv. Lucullus.30 Plants are used 6 weeks after subculture on medium  $A_1$ .- Infection

- Midrib is removed from the leaves.

- Leaves are cut in segments of about 0.25 to 1  $cm^2$  (the edges of the leaves are not wounded, so that only maximum 3 sides of the leaf pieces is wounded).35 - Segments are placed in infection medium  $B_1$ .



(upside down), about 10 segments in a 9 cm Petri dish.

- Segments are then infected with 20  $\mu$ l per Petri dish of a late log culture of the Agrobacterium strain grown in min A medium.

- Petri dishes incubate for 2 days at low light intensity.

- Medium is removed after 2 days and replaced by 20 ml of medium B<sub>1</sub> containing 500 mg/l clarofan.

#### 10 - Selection and shoot induction

- The leaf discs are placed in medium B<sub>2</sub> + 50 or 100 mg/l kanamycin.

- Each 5 days the osmotic pressure of the medium is lowered by decreasing the mannitol concentration, transfers are done consecutively in medium B<sub>3</sub>, B<sub>4</sub>, B<sub>5</sub>, and B<sub>6</sub>.

- After one month calli with meristems are separated from the leaf discs and placed on medium B<sub>7</sub> with 50 or 100 mg/l kanamycin.

- Once little shoots have formed, calli are transferred to elongation medium B<sub>9</sub> with 50 or 100 mg/l kanamycin.

#### - Rooting

- Elongated shoots are separated and transferred to medium B<sub>8</sub> for rooting.

- Plants are propagated on medium A<sub>1</sub>.

### Greenhouse tests for herbicide resistance

#### Material and method

30 In this experiment, two herbicides comprising phosphinotricin as active ingredient, are used.

These compounds are those commercially available under the registered trademarks BASTA<sup>R</sup> and MEIJI HERBIACE<sup>R</sup>.

35 These products are diluted to 2 % with tap water. Spraying is carried out on a square metre area from the

four corners. Temperature of the greenhouse is about 22°C for tobacco, and tomato, and above 10°C to 15°C for potato.

5

### Results

#### - Tobacco spraytest

a) Nicotiana tabacum cv. Petit Havana SR1 plants transformed with the chimeric "sfr" genes as present in pGSFR1161 or pGSFR1281, as well as untransformed control plants (from 10 cm to 50 cm high) are treated with 20 l BASTA<sup>R</sup>/ha. Control SR1 plants die after 6 days, while transformed plants are fully resistant to 20 l BASTA<sup>R</sup>/ha and continue growing undistinguishable from untreated plants. No visible damage is detected, also if the treatment is repeated every two weeks. The treatment has no effect in subsequent flowering. The recommended dose of BASTA<sup>R</sup> herbicide in agriculture is 2.5-7.5 l/ha.

b) A similar experiment is performed using 8 l/ha MEIJI HERBIACE<sup>R</sup>. The transformed plants (the same as above) are fully resistant and continue growing undistinguishable from untreated plants. No visible damage is detectable.

#### - Potato spraytest

Untransformed and transformed potato plants (Solanum tuberosum cv. Berolina) (20 cm high) with the chimeric "sfr" gene as present in pGSFR1161 or pGSFR1281 are treated with 20 l BASTA<sup>R</sup>/ha. Control plants die after 6 days while the transformed plants do not show any visible damage. They grow undistinguishable from untreated plants.

#### - tomato spraytest

Untransformed and transformed tomato plants (lycopersicum esculentum c.v. luculus) (25 cm high) with the chimeric "sfr" gene as present in pGSFR1161 and pGSFR1281 are treated with 20 l BASTA<sup>R</sup>/ha. Control plants

die after six days while transformed plants are fully resistant. They do not show any visible damage and grow undistinguishable from untreated plants.

- 5        - Growth control of phytopathogenic fungi with transformed plants

10        In another set of experiments, potato plants expressing chimeric "sfr" genes as present in pGSFR1161 or pGSFR1281 are grown in a greenhouse compartment at 20°C under high humidity. Plants are inoculated by spraying 1 ml of a suspension of  $10^6$  Phytophthora infestans spores per ml. Plants grow in growth chambers (20°C, 95 % humidity, 4 000 lux) until fungal disease symptoms are visible (one week). One set of the plants are at that moment sprayed with Bialaphos at a dose of 8 l/ha. Two weeks later, 15        untreated plants are completely ingested by the fungus. The growth of the fungus is stopped on the Bialaphos treated plants and no further disease symptoms evolve. The plants are effectively protected by the fungicide action of Bialaphos.

- 20        - Transmission of the PPT resistance through seeds  
      Transformed tobacco plants expressing the chimeric "sfr" gene present in pGSFR1161 and pGSFR1281 are brought to flowering in the greenhouse. They show a normal fertility.

25        About 500 F1 seeds of each plant are sown in soil, F1 designating seeds of the first generation, i.e directly issued from the originally transformed plants. When seedlings are 2-3 cm high, they are sprayed with 8 l BASTA<sup>R</sup>/ha. 7 days later, healthy and damaged plants can be 30        distinguished in a ratio of approximately 3 to 1. this shows that PPT resistance is inherited as a dominant marker encoded by a single locus.

35        10 resistant F1 seedlings are grown to maturity and seeds are harvested. F2 seedlings are grown as described above and tested for PPT-resistance by spraying

BASTA<sup>R</sup> at a dose of 6 l/ha. Some of the F1 plants produce F2 seedlings which are all PPT-resistant showing that these plants are homozygous for the resistance gene. The invention also concerns plant cells and plants non-essentially-biologically-transformed with a GS inhibitor-inactivating-gene according to the invention.

In a preferred embodiment of the invention, plant cells and plants are non-biologically-transformed with the "sfr" gene hereabove described.

Such plant cells and plants possess, stably integrated in their genome, a non-variety-specific character which render them able to produce detectable amounts of phosphinotricin-acetyl transferase.

This character confers to the transformed plant cells and plants a non-variety-specific enzymatic activity capable of inactivating or neutralizing GS inhibitors like Bialaphos and PPT.

Accordingly, plant cells and plants transformed according to the invention are rendered resistant against the herbicidal effects of Bialaphos and related compounds.

Since Bialaphos was first described as a fungicide, transformed plants can also be protected against fungal diseases by spraying with the compound several times.

In a preferred embodiment, Bialaphos or related compounds is applied several times, particularly at time intervals of about 20 to 100 days.

The invention also concerns a new process for selectively protecting a plant species against fungal diseases and selectively destroying weeds in a field comprising the steps of treating a field with an herbicide, wherein the plant species contain in their genome a DNA fragment encoding a protein having an enzymatic activity capable of neutralizing or inactivating GS inhibitors and wherein the used herbicide comprises as

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active ingredient a GS inhibitor.

It comes without saying that the process according to the invention can be employed with the same efficiency, either to only destroy weeds in a field, if plants are not  
5 infected with fungi, either to only stop the development of fungi if the latter appears after destruction of weeds.

In a preferred embodiment of the process according to the invention, plant species are transformed with a DNA  
10 fragment comprising the "sfr" gene as described hereabove, and the used herbicide is PPT or a related compound.

Accordingly, a solution of PPT or related compound is applied over the field, for example by spraying, several times after emergence of the plant species to be  
15 cultivated, until early and late germinating weeds are destroyed.

It is quite evident that before emergence of plant species to be cultivated, the field can be treated with an herbicidal composition to destroy weeds.

On the same hand, fields can be treated even  
20 before the plant species to be cultivated are sowed.

Before emergence of the desired plant species, fields can be treated with any available herbicide, including Bialaphos-type herbicides.

After emergence of the desired plant species,  
25 Bialaphos or related compound is applied several times.

In a preferred embodiment, the herbicide is applied at time intervals of about from 20 to 100 days.

Since plants to be cultivated are transformed in  
30 such a way as to resist to the herbicidal effects of Bialaphos-type herbicides, fields can be treated even after emergence of the cultivated plants.

This is particularly useful to totally destroy early and late germinating weeds, without any effect on  
35 the plants to be produced.

Preferably, Bialaphos or related compound is

5 applied at a dose ranging from about 0.4 to about 1.6 kg/ha, and diluted in a liquid carrier at a concentration such as to enable its application to the field at a rate ranging from about 2 to about 8 l/ha.

There follows examples, given by way of illustration, of some embodiments of the process with different plant species.

- Sugarbeets

10 The North European sugarbeet is planted from March 15 up to April 15, depending upon the weather condition and more precisely on the precipitation and average temperature. the weeds problems are more or less the same in each country and can cause difficulties until the crop  
15 closes its canopy around mid-July.

Weed problems can be separated in three situations :

- early germination of the grassy weeds,
- early germinating broadleaved weeds,
- 20 - late germinating broadleaved weeds.

Up to now, pre-emergence herbicides have been successfully used. Such compounds are for example those commercially available under the registered trademarks : PYRAMIN<sup>R</sup>, GOLTIX<sup>R</sup> and VENZAR<sup>R</sup>. However, the susceptibility  
25 to dry weather conditions of these products as well as the lack of residual activity to control late germinating weeds have led the farmer to use post-emergence products in addition to pre-emergence ones.

30 Table (I) thereafter indicates the active ingredients contained in the herbicidal compositions cited in the following examples.

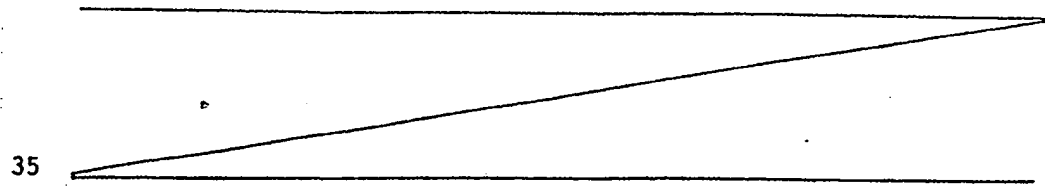


TABLE (I)

	Commercial Name	Active Ingredient	Formulation
5	AVADEX <sup>R</sup>	Diallate	EC 400 g/l
	AVADEX BW <sup>R</sup>	Triallate	EC 400 g/l
	GOLTIX <sup>R</sup>	Metamitron	WP 70 %
	RONEET <sup>R</sup>	Cycloate	EC 718 g/l
10	TRAMAT <sup>R</sup>	Ethofumerate	EC 200 g/l
	FERVINAL <sup>R</sup>	Alloxydime-sodium	SP 75 %
	BASTA <sup>R</sup>	Phosphinotricin	200 g/l
	PYRAMIN FL <sup>R</sup>	Chloridazon	SC 430 g/l

15 According to the invention, post-emergence herbicides consist of Bialaphos or related compounds, which offer a good level of growth control of annual grasses (Bromus, Avena spp., Alopecurus, POA) and broadleaves (Galium, Polygonum, Senecio, Solanum, Mercurialis).

20 Post-emergence herbicides can be applied at different moments of the growth of sugarbeet ; at a cotyledon level, two-leave level or at a four-leave level.

25 Table (II) thereafter represents possible systems of field-treatment, given by way of example.

In those examples, the post-emergence herbicide of the class of Bialaphos used is BASTA<sup>R</sup>, in combination with different pre-emergence herbicides. Concentrations are indicated in l/ha or kg/ha.

30

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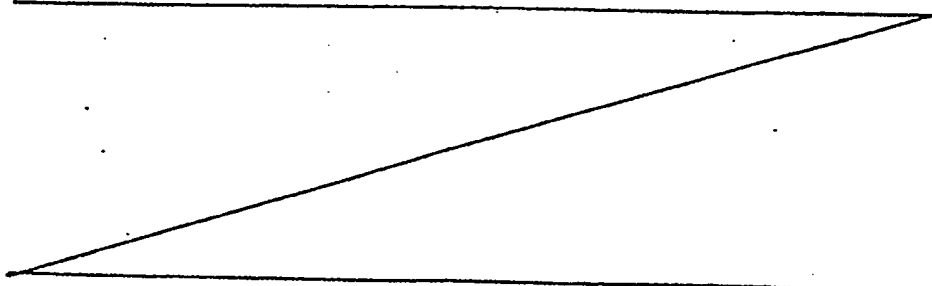


TABLE (II)

POSSIBLE WEEDCONTROL SYSTEMS IN SUGARBEETS, BASED ON THE USE OF BASTA<sup>R</sup>, PROVIDING BEETS ARE MADE RESISTANT AGAINST THE LATTER CHEMICAL (in lt or kg/ha).

	Pre-sowing	Pre-emergence	Cotyledons	Two-leaves	Four leaves
1.	AVADEX <sup>R</sup> 3.5 lt	-	BASTA <sup>R</sup> 3 lt	BASTA <sup>R</sup> /tramat 3 lt 1.5 lt	-
2.	AVADEX <sup>R</sup> 3.5 lt	GOLTIX <sup>R</sup> 4 kg	-	-	-
3	RONEET <sup>R</sup> 4 lt	GOLTIX <sup>R</sup> 5 kg	-	-	-
4.	RONEET <sup>R</sup> 4 lt	GOLTIX <sup>R</sup> 2.5 kg	-	BASTA <sup>R</sup> 3 lt	-
5.	TRAMAT <sup>R</sup> 5 lt	-	-	BASTA <sup>R</sup> 3 lt	BASTA <sup>R</sup> /GOLTIX <sup>R</sup> 2 lt 2 kg
6.	-	GOLTIX <sup>R</sup> 2.5 kg	-	BASTA <sup>R</sup> 3 lt	-
7.	-	-	BASTA <sup>R</sup> /tramat 3 lt 1.7 lt	-	BASTA <sup>R</sup> /GOLTIX <sup>R</sup> 3 lt 2 kg
8.	PYRAMIN <sup>R</sup> 6 lt	-	BASTA <sup>R</sup> 3 lt	Venzar 1 kg	-
9.	-	-	BASTA <sup>R</sup> 3 lt	BASTA <sup>R</sup> /GOLTIX <sup>R</sup> 3 lt 2 kg	-
10.	DIALATE <sup>R</sup> 3.5 lt	PYRAMIN <sup>R</sup> 6 lt	-	BASTA <sup>R</sup> /Metamitron 3 lt 1 kg	-



- Potatoes

Potatoes are grown in Europe on about  $8.10^6$  Ha. The major products used for weed control are Linuron/monolinuron or the compound commercially available under the denomination METRABUZIN.

These products perform well against most weedspecies.

However, weeds such as Galium and Solanum plus late germinating Chenopodium and Polygonum are not always effectively controlled, while control of the annual grasses is also sometime erratic.

Once again, late germinating broadleaved weeds are only controllable by post-emergence applications of herbicides such as BASTA<sup>R</sup>.

Table (III) thereafter represents some examples given by way of example of field-treatment in the case of potatoes.

TABLE (III)

Weeds control systems in potatoes, based on the use of BASTA<sup>R</sup>, providing potatoes are rendered resistant to BASTA<sup>R</sup>.

25

Linuron + monolinuron (375 g + 375 g/ha) prior to emergence

BASTA<sup>R</sup> 3-4 lt/ha after emergence (5-15 cm)

30 BASTA<sup>R</sup>/fluazifop-butyl 3-4 lt/ha + 2 lt/ha after emergence (5-15 cm)

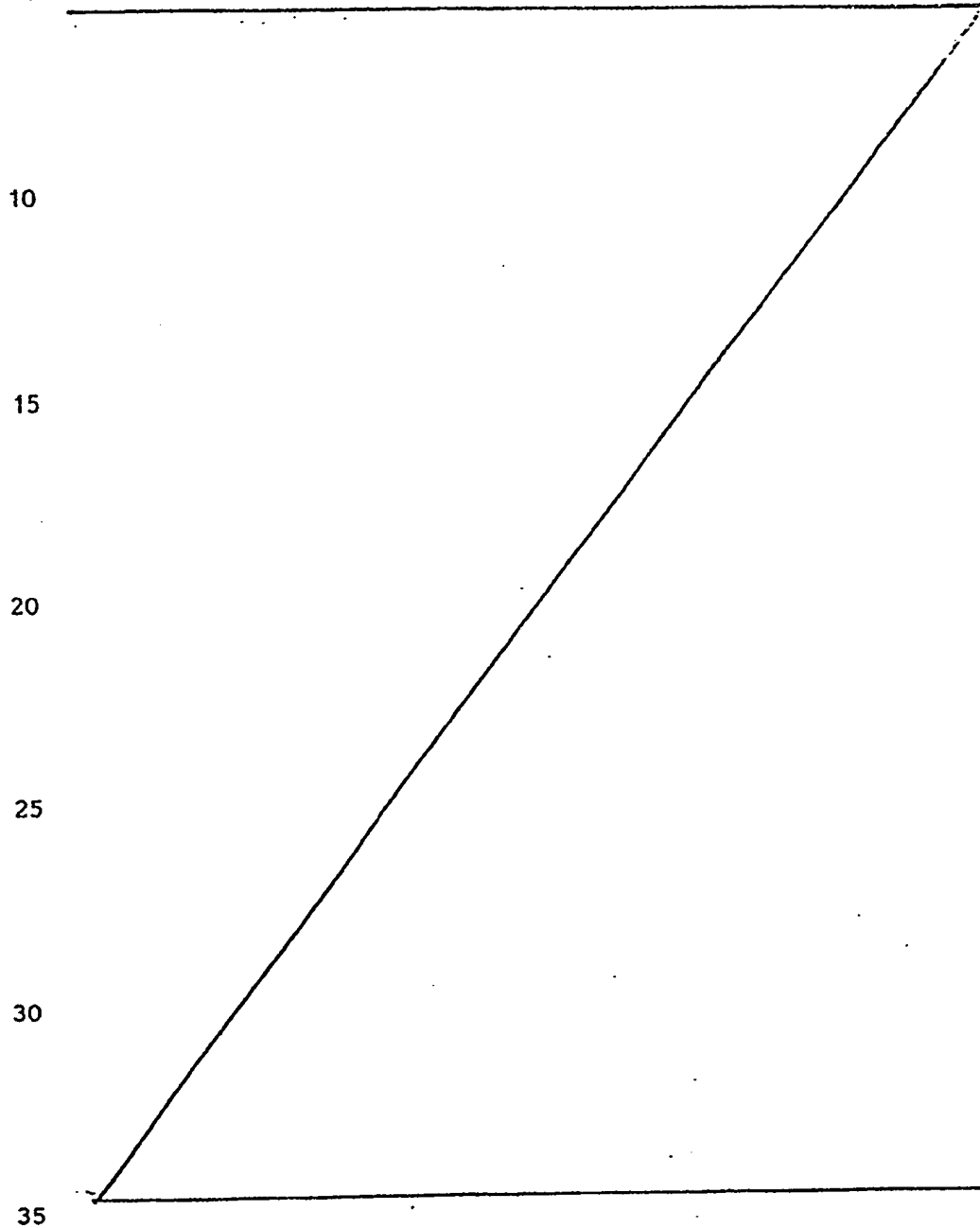
Linuron WP 50 % (AFALON<sup>R</sup>).

Monolinuron WP 47.5 % (ARESSIN<sup>R</sup>)

35 fluazifop-butyl EL 250 g/l (FUSILADE<sup>R</sup>)

The strains pGSJ260 and pBG39 used hereabove have

been deposited on December 12nd, 1985, at the "German Collection of Micro-organisms" (Deutsche Sammlung von Mikroorganismen) at Göttingen, Germany. They received the deposition numbers DSM 3 606 and DSM 3 607 respectively.



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## CLAIMS

1. Process for controlling the action in plant cells and plants of a glutamine synthetase inhibitor when  
5 contacted herewith, which comprises providing said plants with a heterologous DNA including a foreign nucleotide sequence, capable of being expressed in the form of a protein in said plant cells and plants, under conditions such as to cause said heterologous DNA to be integrated  
10 stably through generations in the cells of said plants, and wherein said protein has an enzymatic activity capable of causing inactivation or neutralization of said glutamine synthetase inhibitor.

2. Process according to claim 1, wherein the  
15 heterologous DNA fragment comprises a foreign nucleotide sequence coding for a polypeptide having a PPT acetyl transferase activity.

3. Process according to the claims 1 to 2, wherein the heterologous DNA fragment belongs to the genome of a  
20 Streptomyces.

4. Process according to any of the claims 1 to 3, wherein the heterologous DNA fragment comprises foreign

25

30

35



in which X represents MET or VAL, which part of said protein is of sufficient length to possess said enzymatic activity and, when said DNA fragment is expressed in plant cells, to protect the latter against the herbicidal activity of glutamine synthetase inhibitors.

5. Process according to any of the claims 1 to 4, wherein the heterologous DNA fragment comprises the following nucleotide sequence :

GTG AGC CCA GAA  
 183  
 CGA CGC CCG GCC GAC ATC CGC CGT GCC ACC GAG GCG GAC ATG CCG  
 228  
 GCG GTC TGC ACC ATC GTC AAC CAC TAC ATC GAG ACA AGC ACG GTC  
 273  
 AAC TTC CGT ACC GAG CCG CAG GAA CCG CAG GAG TGG ACG GAC GAC  
 318  
 CTC GTC CGT CTG CCG GAG CCG TAT CCC TGG CTC GTC GCC GAG GTG  
 363  
 GAC GGC GAG GTC GCC GGC ATC GCC TAC GCG GGC CCC TGG AAG GCA  
 408  
 CGC AAC GCC TAC GAC TGG ACG GCC GAG TCG ACC GTG TAC GTC TCC  
 453  
 CCC CGC CAC CAG CCG ACG GGA CTG GGC TCC ACG CTC TAC ACC CAC  
 498  
 CTG CTG AAG TCC CTG GAG GCA CAG GGC TTC AAG AGC GTG GTC GCT  
 543  
 GTC ATC GGG CTG CCC AAC GAC CCG AGC GTG CCG ATG CAC GAG GCG  
 588  
 CTC GGA TAT GCC CCC CGC GGC ATG CTG CCG GCG GCC GGC TTC AAG  
 633  
 CAC GGG AAC TGG CAT GAC GTG GGT TTC TGG CAG CTG GAC TTC AGC  
 678  
 CTG CCG GTA CCG CCC CGT CCG GTC CTG CCC GTC ACC GAG ATC  
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6. Process according to any of claims 1 to 5, wherein the initiation codon ATG is substituted for the initiation codon GTG of the sequence defined in claim 5.

5 7. In a process for producing plants and reproduction material of said plants including a heterologous genetic material stably integrated therein and capable of being expressed in said plants or reproduction material in the form of a protein capable of  
10 inactivating or neutralizing the activity of a glutamine synthetase inhibitor, comprising the non biological steps of producing plants cells or plant tissue including said heterologous genetic material from starting plant cells or plant tissue not able to express that inhibiting or  
15 neutralizing activity, regenerating plants or reproduction material of said plants or both from said plant cells or plant tissue including said genetic material and, optionally, biologically replicating said last mentioned plants or reproduction material or both, wherein said  
20 steps of producing said plant cells or plant tissue including said heterologous genetic material comprises transforming said starting plant cells or plant tissue with a DNA recombinant containing a nucleotide sequence encoding said protein as well as the regulatory elements  
25 selected among those which are capable of enabling the expression of said nucleotide sequence in said plant cells or plant tissue and to cause the stable integration of said nucleotide sequence in said plant cells and tissue, as well as in the plant and reproduction material  
30 processed therefrom throughout generations.

8. The process according to claim 7, wherein starting cells are transformed with a recombinant DNA which contains the fragment of any one of the claims 1 to 6 recombined with heterologous DNA.

35 9. The process according to claim 8, wherein the

recombinant DNA is a vector suitable for the transformation of a cellular host and, when introduced therein, for allowing for the neutralization or inhibition of glutamine synthetase inhibitors.

5 10. The process according to claim 10, wherein the vector comprises said recombinant DNA under the control of replicon elements suitable for the transformation of bacteria, particularly E.coli.

10 11. The process of claim 9, wherein the vector comprises said recombinant DNA fragment under the control of a plant promotor region, which vector further comprises regulation elements allowing for the expression of said DNA fragments in plant cells, when the latter are later transformed with said vector.

15 12. The process of any of claims 8 to 11, which confers resistance against herbicidal effects of Bialaphos, PPT or related derivatives to the transformed plant cells.

20 13. The process of claim 11 or 12, wherein the vector comprises a nucleotide sequence encoding a transit peptide intercalated between said plant promotor region and said recombinant DNA fragment.

25 14. The process of claim 13, wherein the transit peptide is selected from ribulose-1,5 biphosphate carboxylase and chlorophyl a/b binding proteins.

15. The process of any of claims 9 to 14, wherein said vector is a Ti plasmid.

30 16. Plant cells, non biologically transformed, wherein they possess, stably integrated in their genome, a DNA fragment encoding a protein having a non-variety-specific enzymatic activity capable of neutralizing or inactivating glutamine synthetase inhibitors.

35 17. Plant cells according to claim 16, wherein they are capable of generating a plant capable of producing seeds, said seeds having a non-variety-specific

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enzymatic activity capable of inactivating or neutralizing glutamine synthetase inhibitors.

18. Plant cells according to claim 16 or 17,  
5 wherein they are transformed by the process of any of claims 7 to 15.

19. Plant cells according to any of claims 16 to 18, which \_\_\_\_\_ produce detectable amounts of phosphotricin acetyl transferase.

10 20. Seeds, non biologically transformed, which possess, stably integrated in their genome, a DNA fragment encoding a protein having a non-variety-specific enzymatic activity capable of inactivating or neutralizing glutamine synthetase inhibitors.

15 21. Seeds according to claim 20, which \_\_\_\_\_ are capable of germinating into a plant capable of producing seeds having a non-variety-specific enzymatic activity capable of inactivating or neutralizing glutamine synthetase inhibitors.

20 22. Seeds according to \_\_\_\_\_ claim 20 or 21, which are transformed by the process of any one of the claims 7 to 15.

25 23. Plants, non biologically transformed, which possess, stably integrated in their genome, a DNA fragment encoding a protein having a non-variety-specific enzymatic activity capable of neutralizing or inactivating glutamine synthetase inhibitors.

30 24. Plants according to claim 23, which are capable of producing seeds having a non-variety-specific enzymatic activity capable of inactivating or neutralizing glutamine synthetase inhibitors.

25. Plants according to claim 23 or 24, which are transformed by the process of any of \_\_\_\_\_ claims 7 to 15.

35 26. Process for protecting a plant species and selectively destroying weeds in a field comprising the steps of treating a field with an herbicide, wherein the

plant species contain in their genome a DNA fragment encoding a protein having an enzymatic activity capable of neutralizing or inactivating glutamine synthetase inhibitors, and wherein the used herbicide is a glutamine synthetase inhibitor.

27. Process according to claim 26, wherein the plant species contain a DNA fragment according to any one of claims 1 to 6.

28. Process according to claim 26 or 27, wherein the plant species are transformed according to the process of any one of claims 7 to 15.

29. Process according to any one of claims 26 to 28, wherein a solution of a glutamine synthetase inhibitor is applied on the field after emergence of the cultivated plant species, several times, particularly at time intervals of about 20 to 100 days, until early and late germinating weeds are destroyed.

30. Process according to any of claims 26 to 29, wherein glutamine synthetase inhibitors comprise Bialaphos, phosphinotricin and related compounds.

31. Process for selectively protecting a plant species in a field against fungal diseases comprising the steps of treating a field with an herbicide, wherein the plant species contain in their genome a DNA fragment encoding a protein having an enzymatic activity capable of neutralizing an inactivating glutamine synthetase inhibitors and wherein the used herbicide is a glutamine synthetase inhibitor.

32. Process according to claim 31, wherein the plant species contain a DNA fragment according to any of claims 1 to 6.

33. Process according to claim 31 or 32, wherein the plant species are transformed according to the process of any one of claims 7 to 15.

34. Process according to any one of claims 31 to

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33, wherein a solution of a glutamine synthetase inhibitor is applied on the field after emergence of the cultivated plant species, several times, particularly at \_\_\_\_\_ time intervals of about 20 to 100 days until fungi are destroyed.

35. Process according to any one of claims 31 to 34, wherein glutamine synthetase inhibitor is selected from a group which comprises Bialaphos, phosphinotricin and related compounds.

36. Process according to claim 30 or 35, wherein Bialaphos, PPT or related compound is applied at a dose ranging from about 0.4 to about 1.6 kg/ha.

37. Process according to claim 36, wherein Bialaphos, PPT or related compound is diluted in a liquid carrier at a concentration such as to enable its application to the field at a rate ranging from about 2 l/ha to about 8 l/ha.

38. Process according to any of claim 36 or 37, wherein selectively protected plant species comprise, sugar-beet, rice, potato, tomato, maize, tobacco.

39. Vector which contains a DNA fragment encoding a polypeptide having an enzymatic activity capable of causing inactivation or neutralization of glutamine synthetase inhibitors.

40. Vector according to claim 39, which is suitable for the transformation of plant cells and plants.

41. Vector according to claim 39 or 40, which contains a DNA fragment belonging to the genome of a Streptomyces.

42. Vector according to any of \_\_\_\_\_ claims 39 to 41, which comprises the following nucleotide sequence :

GTG AGC CCA GAA

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5 CGA CGC CCG GCC GAC ATC CGC CGT GCC ACC GAG GCG GAC ATG CCG  
 228  
 GCG GTC TGC ACC ATC GTC AAC CAC TAC ATC GAG ACA AGC ACG GTC  
 273  
 AAC TTC CGT ACC GAG CCG CAG GAA CCG CAG GAG TGG ACG GAC GAC  
 318  
 10 CTC GTC CGT CTG CCG GAG CGC TAT CCC TGG CTC GTC GCC GAG GTG  
 363  
 GAC GGC GAG GTC GCC GGC ATC GCC TAC GCG GGC CCC TGG AAG GCA  
 408  
 15 CCG AAC GCC TAC GAC TGG ACG GCC GAG TCG ACC GTG TAC GTC TCC  
 453  
 CCC CGC CAC CAG CCG ACG GGA CTG GGC TCC ACG CTC TAC ACC CAC  
 498  
 CTG CTG AAG TCC CTG GAG GCA CAG GGC TTC AAG AGC GTG GTC GCT  
 543  
 20 GTC ATC GGG CTG CCC AAC GAC CCG AGC GTG CGC ATG CAC GAG GCG  
 588  
 CTC GGA TAT GCC CCC CGC GGC ATG CTG CCG GCG GCC GGC TTC AAG  
 633  
 25 CAC GGG AAC TGG CAT GAC GTG GGT TTC TGG CAG CTG GAC TTC AGC  
 678  
 CTG CCG GTA CCG CCC CGT CCG GTC CTG CCC GTC ACC GAG ATC  
 723

30

35

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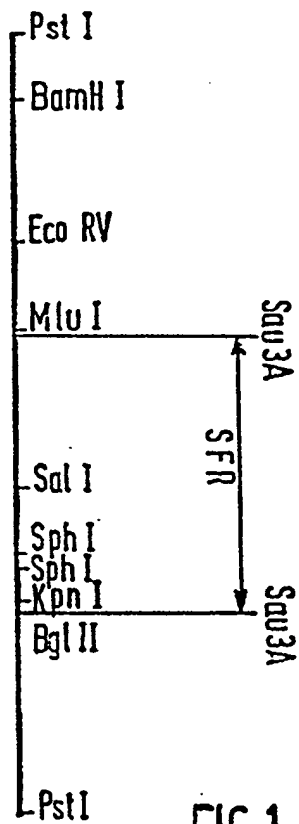


FIG.1

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CCC GCT CAA GCT CGC TGT CAT TTT CGA GAC GCC ATC TTT GGA AGC  
 GGT GGC CGA ATC CGT ACT GCG CGG ACT CGA CGA CGC GTA AAA CGA  
 TCG ACC ACG TAC ACG AGT CCG GAC ACG GGG CGA GGA GGC CCG GTT  
 CCG GCA CCG AGG AAG ACC GAA GGA AGA CCA CAC GTG AGC CCA GAA  
 CCG CGC CCG GCC GAC ATC CCG CGT GCC ACC GAG GCG GAC ATG CCG  
 FokI  
 GCG GTC TGC ACC ATC GTC AAC CAC TAC ATC GAG ACA AGC ACG GTC  
 AAC TTC CGT ACC GAG CCG CAG GAA CCG CAG GAG TGG ACG GAC GAC  
 CTC GTC CGT CTG CCG GAG CCG TAT CCC TGG CTC GTC GCC GAG GTG  
 GAC GCG GAG GTC GCC GGC ATC GCC TAC GCG GGC CCC TGG AAG GCA  
 CCG AAC GCC TAC GAC TGG ACG GCC GAG TCG ACC GTG TAC GTC TCC  
 CCC CGC CAC CAG CCG ACG GGA CTG GGC TCC ACG CTC TAC ACC CAC  
 CTG CTG AAG TCC CTG GAG GCA CAG GGC TTC AAG AGC GTG GTC GCT  
 GTC ATC GGG CTG CCC AAC GAC CCG AGC GTG CCG ATG CAC GAG GCG  
 CTC GGA TAT GCC CCC CCG GGC ATG CTG CCG GCG GCC GGC TTC AAG  
 CAC GGG AAC TGG CAT GAC GTG GGT TTC TGG CAG CTG GAC TTC AGC  
 CTG CCG GTA CCG CCC CGT CCG GTC CTG CCC GTC ACC GAG ATC TGA  
 BglII  
 ACG GAG TGC GCG TGG GCA TCG CCC GAG TTG GAG CTG GTA CCG GAA  
 CTC ATC GAA CTC AAC TGG CAT ACC CCG AAT GGT GAG GTG GAA CCG  
 CCG CCG ATC GCG TAC GAC CGT GCC CAG G

FIG.2

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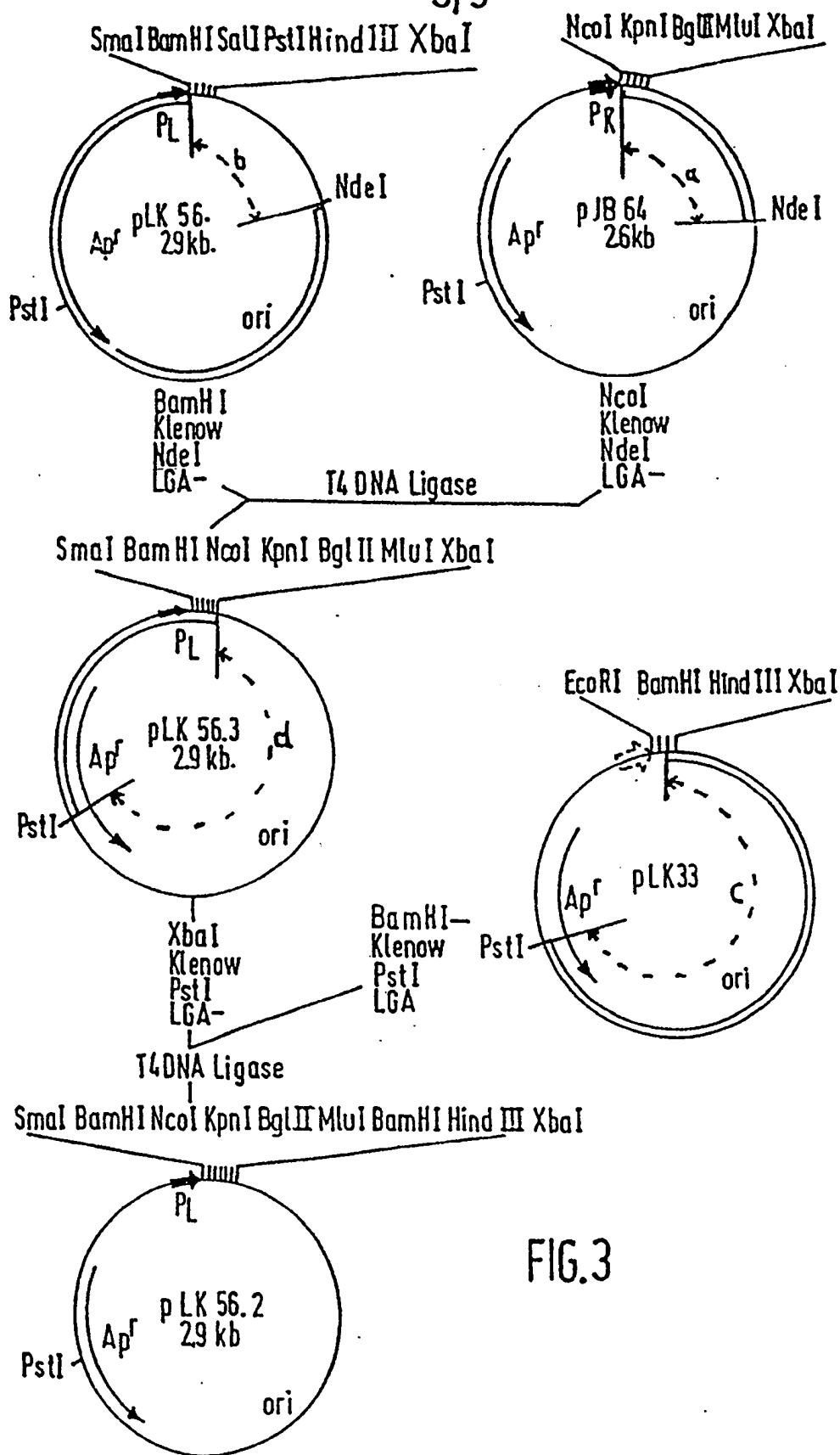


FIG.3

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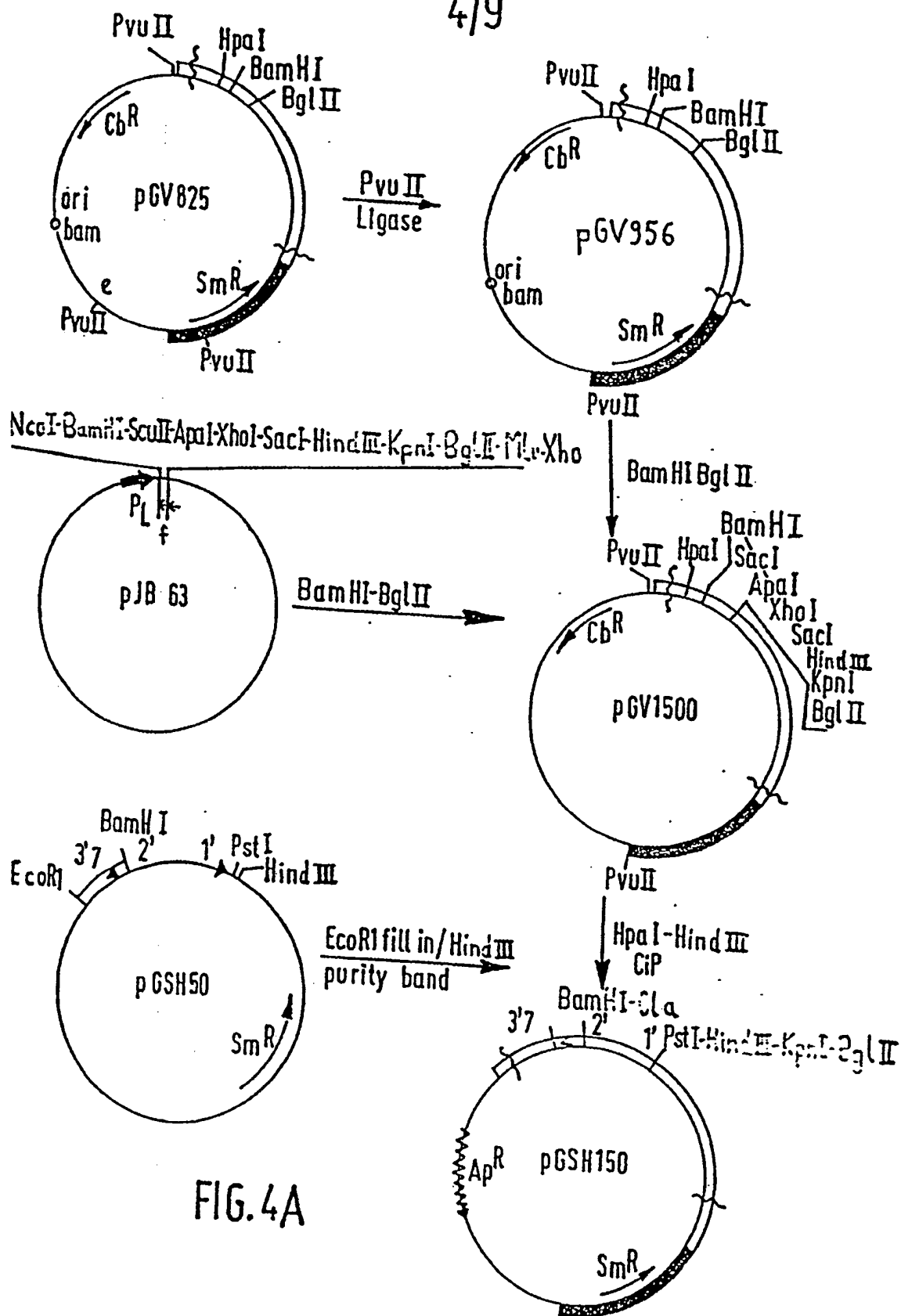
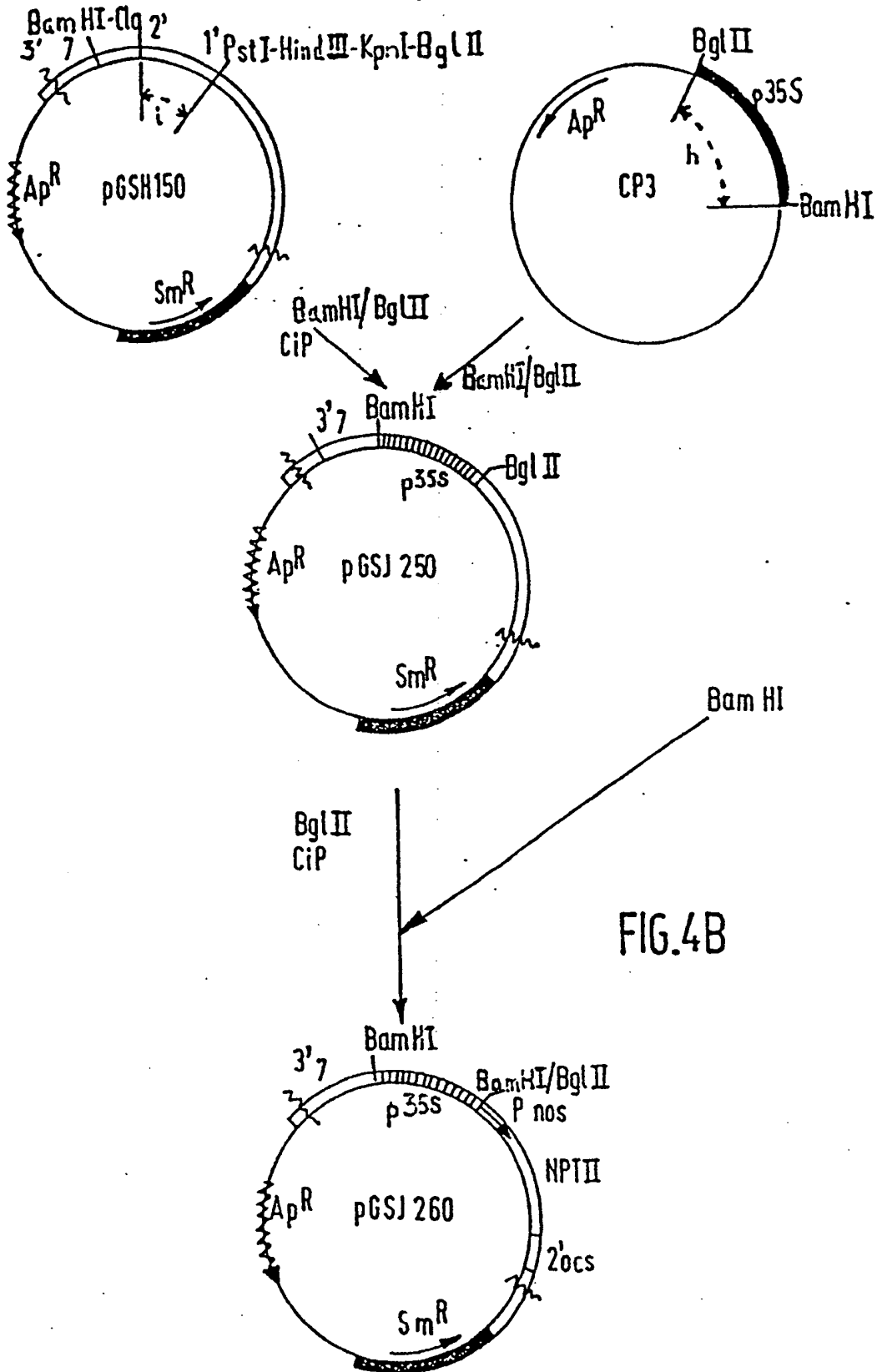


FIG. 4A



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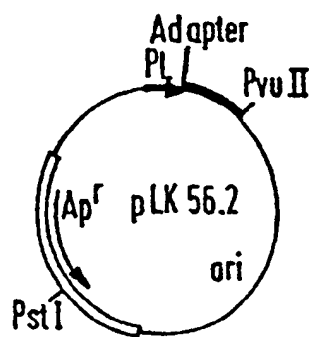


FIG.5A

NcoI / Bgl II

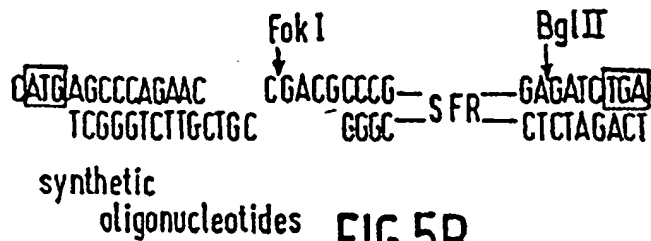


FIG.5B

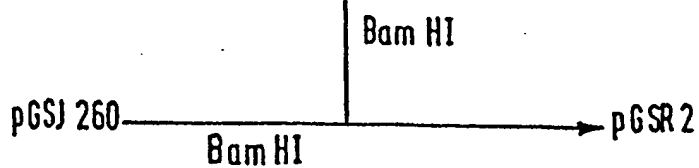
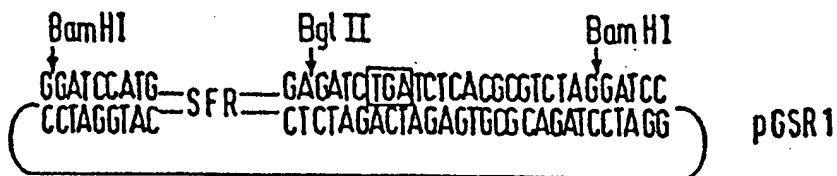
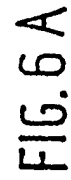
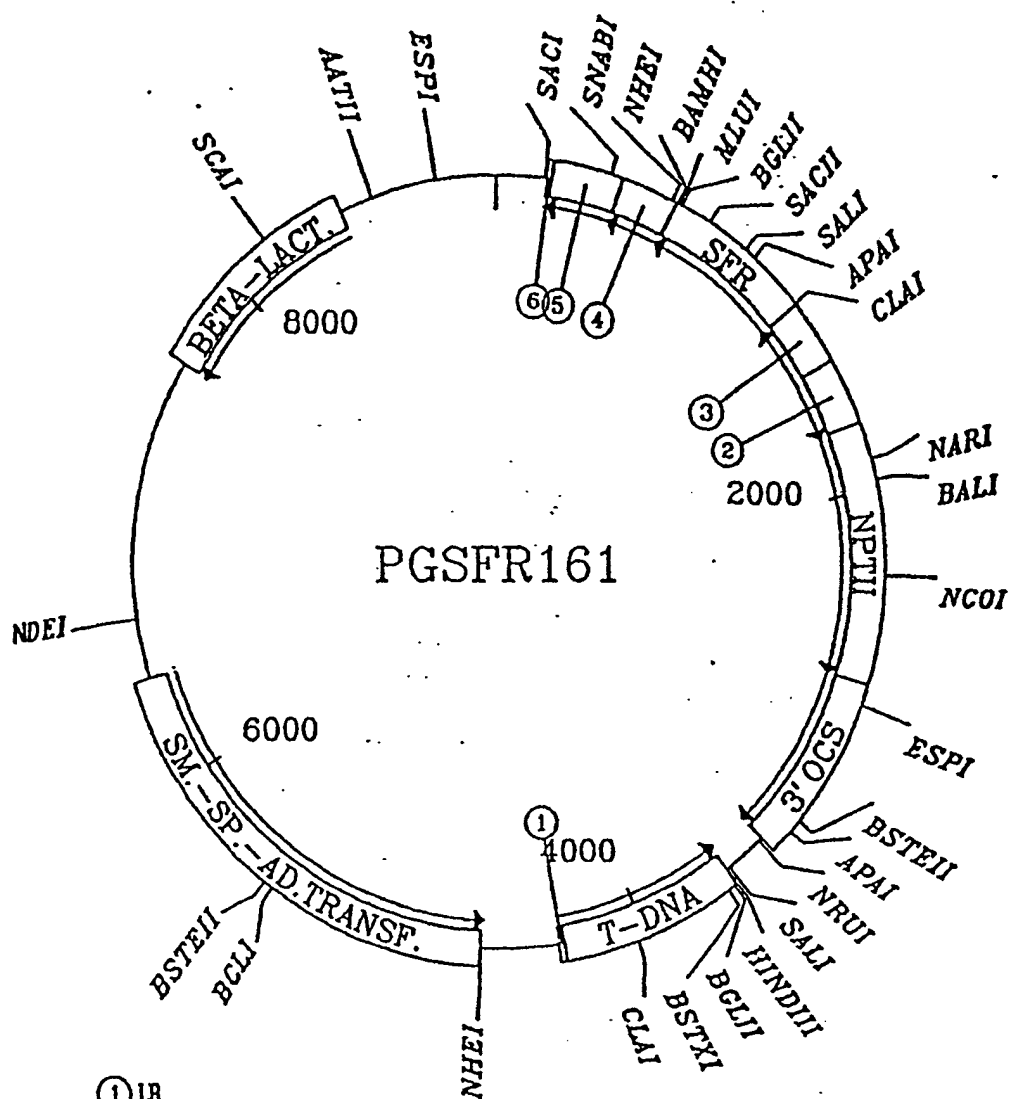


FIG.5C





- ① LB
- ② PTR1
- ③ PTR2
- ④ 3' END 7
- ⑤ T-DNA
- ⑥ R.B.

FIG. 6B

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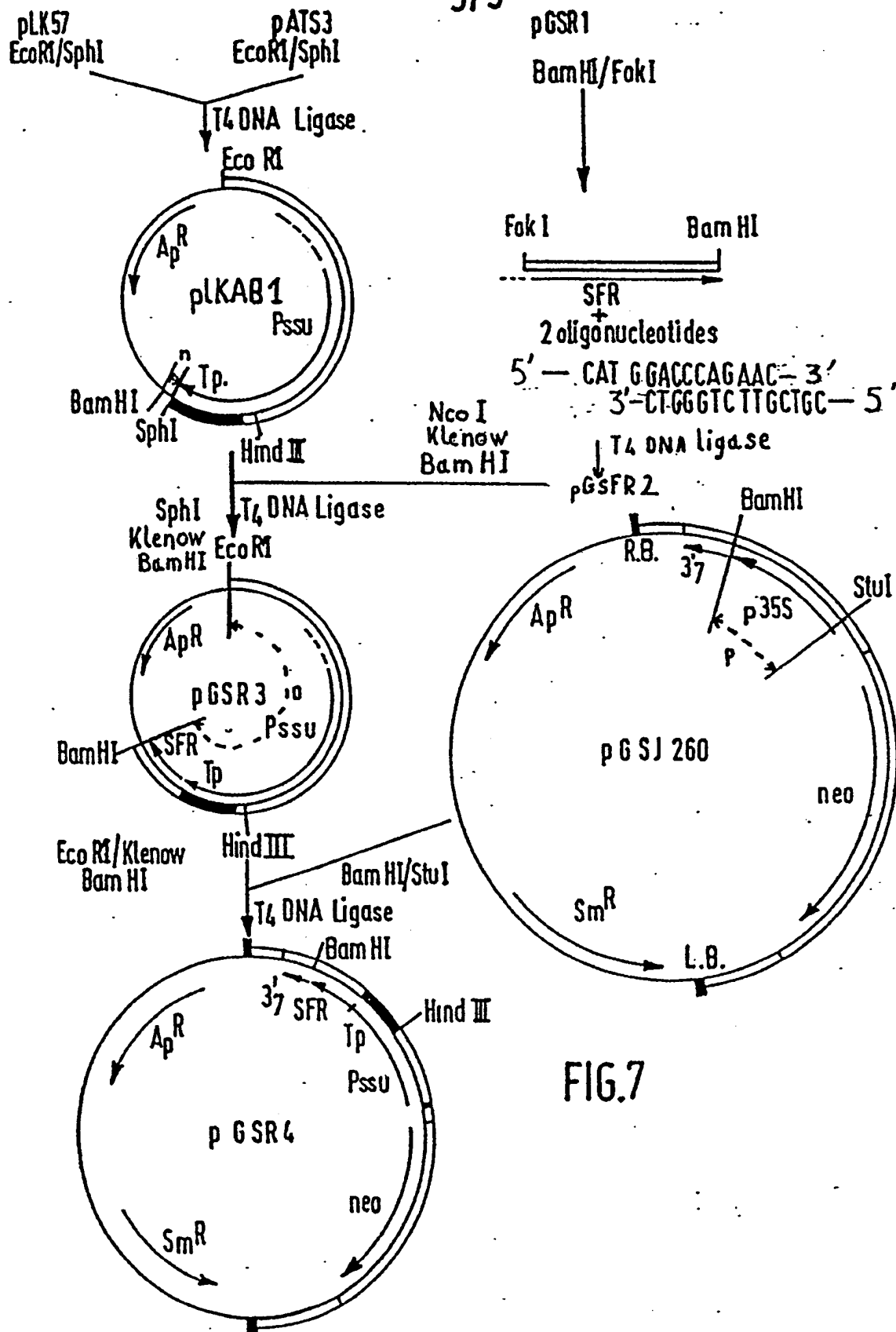


FIG.7



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P,Y	Idem	1-38, 40	
A	EMBO JOURNAL, vol. 4, 1985, pages 25-32, IRL Press Ltd., Oxford, GB; P.H. SCHREIER et al.: "The use of nuclear-encoded sequences to direct the light-regulated synthesis and transport of a foreign protein into plant chloroplasts" * Page 30, right-hand column - page 31 *	11-15	TECHNICAL FIELDS SEARCHED (Int. Cl.4)
A	PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF USA, vol. 81, May 1984, pages 2960-2964; A.R. CASHMORE: "Structure and expression of a pea nuclear gene encoding a chlorophyll a/b-binding polypeptide" * Page 2964 *	11-15	
The present search report has been drawn up for all claims			
Place of search <b>THE HAGUE</b>		Date of completion of the search <b>04-07-1987</b>	Examiner <b>MADDOX A.D.</b>
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DOCUMENTS CONSIDERED TO BE RELEVANT			Page 2
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Y	WO-A-8 402 913 (MONSANTO)  * Page 4, lines 17-28; page 5, lines 9-20; claims 17-19; page 41; figures 18,21,27,37 *	6,13-15	
Y	--- GENE, vol. 33, no. 2, 1985, pages 197-206, Elsevier Science Publishers; J. VARA et al.: "Cloning and expression of a puromycin N-acetyl transferase gene from Streptomyces alboniger in Streptomyces lividans and Escherichia coli" * Summary; page 204, right-hand column, paragraph 1 *	10	
Y	--- WO-A-8 602 097 (GENERAL HOSPITAL CORP.) * Page 13, last paragraph - page 14, paragraph 1; page 26; claim 23 *	26-38	
Y	--- "The Pesticide Manual", page 302, 7th edition, editor C.R. WORTHING, The British Crop Protection Council * Page 302, entry no. 6930 *	26-38	TECHNICAL FIELDS SEARCHED (Int. Cl.4)
The present search report has been drawn up for all claims			
Place of search THE HAGUE		Date of completion of the search 04-07-1987	Examiner MADDOX A.D.
<p><b>CATEGORY OF CITED DOCUMENTS</b></p> <p>X : particularly relevant if taken alone Y : particularly relevant if combined with another document of the same category A : technological background O : non-written disclosure P : intermediate document</p> <p>T : theory or principle underlying the invention E : earlier patent document, but published on, or after the filing date D : document cited in the application L : document cited for other reasons &amp; : member of the same patent family, corresponding document</p>			



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Application number

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Y	---	1+38, 40-42	
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Y	JOURNAL OF MOLECULAR AND APPLIED GENETICS, vol. 2, no. 6, 1984, pages 621-635, Raven Press, New York, US; G. DONN et al.: "Herbicide-resistant alfalfa cells: an example of gene amplification in plants" * Page 625, right-hand column, last paragraph *	1-42	
Y	NATURE, vol. 317, 24th October 1985, page 668; R. SHIELDS: "Engineering herbicide resistance" * Whole article *	1-15	
The present search report has been drawn up for all claims			
Place of search THE HAGUE		Date of completion of the search 04-07-1987	Examiner MADDOX A.D.
<p><b>CATEGORY OF CITED DOCUMENTS</b></p> <p>X : particularly relevant if taken alone Y : particularly relevant if combined with another document of the same category A : technological background O : non-written disclosure P : intermediate document</p> <p>T : theory or principle underlying the invention E : earlier patent document, but published on, or after the filing date D : document cited in the application L : document cited for other reasons</p> <p>&amp; : member of the same patent family, corresponding document</p>			





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A	--- NUCLEIC ACIDS RESEARCH, vol. 14, no. 4, February 1986, pages 1565-1581, IRL Press Ltd., Oxford, GB; M. ZALACAIN et al.: "Nucleotide sequence of the hygromycin B phosphotransferase gene from Streptomyces hygroscopicus" * Page 1573, paragraph 1; page 1577, paragraph 2 * --- -/-	6	
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A	CHIMICAL ABSTRACTS, vol. 104, no. 9, 3rd March 1986, page 311, abstract no. 64619g, Columbus, Ohio, US; M.C. ERICKSON et al.: "Purification and properties of glutamine synthetase from spinach leaves", & PLANT. PHYSIOL. 1985, 79(4), 923-7		
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